10/634,477

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L2
             52 S E3-E7, E9-E12
                E LEHMANN P/AU
            267 S E3-E6, E11-E14
L3
                E OREDDIGER R/AU
                E ROEDDIGER R/AU
L4
              9 S E3, E4
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              2 S E4
L5
                E RODIGER R/AU
              1 S E4
L6
                E RODDIGER R/AU
L7
              2 S E4
                E WALTER MATSUI/AU
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L9
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                SEL RN L1
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L10
              7 S E1-E7
              6 S L10 AND ERYTHROPOIETIN
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L11
L12
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L13
           1792 S E3
L14
           1792 S L11, L13
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              1 S E3
L15
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E FE/MF L16 30 S E3 NOT MASS

L17 30 S L15, L16

#### FILE 'HCAPLUS' ENTERED AT 14:24:56 ON 23 AUG 2005 L18 9810 S L14

L19 11804 S ?ERYTHROPOIETIN? L20

129 S DARBEPOETIN? (S) (ALPHA OR ALFA)

L21 135 S ?DARBEPOETIN? L22 6067 S EPO OR EPREX

L23 298 S EPOETIN? (S) (ALFA OR ALPHA)

L24 100 S EPOETIN? (S) BETA

L25 458 S EPOETIN L26 42 S ARANESP L27 14463 S L18-L26 L28 655 S L27 AND L17 L29 1236 S L27 AND (FE OR IRON) L30 1243 S L28, L29 E HEART DISEASE/CT E E4+ALL

E E2+ALL 86736 S E7+OLD, NT

L31 L32 29 S L30 AND L31

L33 0 S E90+OLD, NT AND L30

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L34
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L35
L36
              1 S L32, L34 AND L1-L9
L37
              3 S L35 AND ?CONJUGAT?
L38
              2 S L37 NOT 3/SC
L39
              2 S L36, L38
L40
             33 S L35 NOT L36-L39
                SEL DN AN 6-9 13-15 19-27
L41
             16 S L40 AND E1-E48
             18 S L39, L41
L42
            597 S ?RHUEPO?
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L44
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L45
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                E E3+ALL
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L48
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L55
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L63
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L65
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L67
             11 S L14 AND S/ELS
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L70
           1808 S L63-L65,L69
L71
           1808 S L70 OR L70
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SET SMARTSELECT OFF

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L78
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L81
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          15844 S L86 FUL
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            86 S L90 AND C2H4O
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L93
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L94
L95
             35 S L94 NOT P/ELS
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L97
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L98
            110 S L98 AND (PY<=2002 OR PRY<=2002 OR AY<=2002)
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L100
             18 S L99 AND L51 AND L52
             0 S L100 AND L31
L101
             6 S L98 AND L31
L102
             4 S L99 AND L31
L103
L104
             6 S L102,L103
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L105
                SEL DN AN 3 7 9 12 13 15 16 17
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L106
               SEL DN AN L48 1 4
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L107
L108
             27 S L49, L106, L107
                SEL HIT RN
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             15 S L109 AND L14
L110
              3 S L109 AND L17
L111
L112
              1 S L109 AND L59
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jan delaval - 23 august 2005

NODE ATTRIBUTES: DEFAULT MLEVEL IS ATOM DEFAULT ECLEVEL IS LIMITED

GRAPH ATTRIBUTES: RING(S) ARE ISOLATED OR EMBEDDED NUMBER OF NODES IS 11

STEREO ATTRIBUTES: NONE

L90 15844 SEA FILE=REGISTRY SSS FUL L86

L91 86 SEA FILE=REGISTRY ABB=ON PLU=ON L90 AND C2H4O

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-3/BI OR 725273-90-5/BI OR 88504-24-9/BI)

=> fil hcaplus
FILE 'HCAPLUS' ENTERED AT 15:25:04 ON 23 AUG 2005
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FILE COVERS 1907 - 23 Aug 2005 VOL 143 ISS 9 FILE LAST UPDATED: 22 Aug 2005 (20050822/ED)

New CAS Information Use Policies, enter HELP USAGETERMS for details.

This file contains CAS Registry Numbers for easy and accurate substance identification.

=> d l108 bib abs hitstr retable tot

L108 ANSWER 1 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN

AN 2004:996201 HCAPLUS

DN 141:422003

TI Cell-free oligosaccharide remodeling and glycoPEGylation methods and the

```
proteins/peptides produced
     De Frees, Shawn; Zopf, David; Bayer, Robert; Bowe, Caryn; Hakes, David;
IN
PA
     Neose Technologies, Inc., USA
so
     PCT Int. Appl., 1024 pp.
     CODEN: PIXXD2
DT
     Patent
LA
     English
FAN.CNT 15
     PATENT NO.
                          KIND
                                  DATE
                                              APPLICATION NO.
                                                                       DATE
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     US 2003-360779
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                                  20030219
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US 2003-448381P P 20030219

AB The invention includes methods and compns. for remodeling a peptide mol., including the addition or deletion of one or more glycosyl groups to a peptide, and/or the addition of a modifying group to a peptide. In vitro methods for addition and/or deletion of sugars to or from a glypeptide mol. are carried out in a manner as to provide a peptide mol. having a specific customized or desired glycosylation pattern, preferably including the addition of a modified sugar. The peptide is enzymically treated in vitro by the systematic addition of the appropriate enzymes and substrates. A key feature of the invention therefore is to take a peptide produced by any cell type and generate a core glycan structure on the peptide, following which the glycan structure is then remodeled in vitro to generate a peptide having a glycosylation pattern suitable for therapeutic use in a mammal. The blood-circulation half-life of the selected peptide is extended by conjugating the peptide to a synthetic or natural polymer of a size sufficient to retard the filtration of the protein by the glomerulus, as illustrated by conjugating erythropoietin to albumin via a polyethylene glycol (PEG) linker using a combination of chemical and enzymic modifications.

IT 11096-26-7P, Erythropoietin

RL: BPN (Biosynthetic preparation); BSU (Biological study, unclassified); PAC (Pharmacological activity); PKT (Pharmacokinetics); THU (Therapeutic use); BIOL (Biological study); PREP (Preparation); USES (Uses) (cell-free oligosaccharide remodeling and glycoPEGylation methods and

the proteins/peptides produced)

RN 11096-26-7 HCAPLUS

CN Erythropoietin (9CI) (CA INDEX NAME)

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

IT 25322-68-3, Poly(ethylene glycol)

RL: BSU (Biological study, unclassified); RCT (Reactant); THU (Therapeutic use); BIOL (Biological study); RACT (Reactant or reagent); USES (Uses) (cell-free oligosaccharide remodeling and glycoPEGylation methods and the proteins/peptides produced)

RN 25322-68-3 HCAPLUS

CN Poly(oxy-1,2-ethanediyl),  $\alpha$ -hydro- $\omega$ -hydroxy- (9CI) (CA INDEX NAME)

$$HO \longrightarrow CH_2 - CH_2 - O \longrightarrow H$$

L108 ANSWER 2 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN

AN 2004:467755 HCAPLUS

DN 141:34188

TI Methods for the use of **erythropoietin** and its derivatives for the treatment of heart diseases

IN Lehmann, Paul; Roeddiger, Ralf; Walter-Matsui, Ruth

PA F. Hoffmann-La Roche A.-G., Switz.

SO PCT Int. Appl., 31 pp.

CODEN: PIXXD2

DT Patent

LA English

FAN.CNT 1

PATENT NO. KIND DATE

APPLICATION NO.

DATE

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      WO 2004047858
                                          20040610
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                                                                                        20031117 <--
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RW: BW, GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM, AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HU, IE, IT, LU, MC, NL, PT, RO, SE, SI, SK, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG
      US 2004209802
                                 A1
                                          20041021
                                                       US 2003-706701
                                                                                       20031112 <--
PRAI EP 2002-26342
                                 Α
                                          20021122
                                                      <--
      The present invention relates to the use of erythropoietin for
      the treatment of disturbances of iron distribution in heart
      diseases.
IT
      702719-61-7, Erythropoietin (human) 702719-62-8
       , Erythropoietin (human)
      RL: PAC (Pharmacological activity); PRP (Properties); THU (Therapeutic
      use); BIOL (Biological study); USES (Uses)
           (amino acid sequence; methods for use of erythropoietin (
          EPO) and its derivs. for treatment of heart diseases)
RN
      702719-61-7 HCAPLUS
      Erythropoietin (human) (9CI) (CA INDEX NAME)
CN
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
RN
      702719-62-8 HCAPLUS
      Erythropoietin (human) (9CI) (CA INDEX NAME)
CN
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
      7439-89-6, Iron, biological studies
TT
      RL: BSU (Biological study, unclassified); BIOL (Biological study)
           (disturbances in cardiac distribution; methods for use of
           erythropoietin (EPO) and its derivs. for treatment of
          heart diseases)
RN
      7439-89-6 HCAPLUS
CN
      Iron (7CI, 8CI, 9CI) (CA INDEX NAME)
Fe
IT
      11096-26-7, Erythropoietin
      RL: PAC (Pharmacological activity); PRP (Properties); THU (Therapeutic
      use); BIOL (Biological study); USES (Uses)
           (methods for use of erythropoietin (EPO) and its
          derivs. for treatment of heart diseases)
RN
      11096-26-7 HCAPLUS
CN
      Erythropoietin (9CI)
                                  (CA INDEX NAME)
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
IT
      11096-26-7D, Erythropoietin, conjugates and
      derivs. 113427-24-0, Epoetin alfa
      122312-54-3, Epoetin beta 209810-58-2
       , Darbepoetin alfa
      RL: PAC (Pharmacological activity); THU (Therapeutic use); BIOL
       (Biological study); USES (Uses)
           (methods for use of erythropoietin (EPO) and its
          derivs. for treatment of heart diseases)
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RN
     11096-26-7 HCAPLUS
CN
     Erythropoietin (9CI)
                         (CA INDEX NAME)
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*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
     122312-54-3 HCAPLUS
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     Erythropoietin [30-asparagine, 32-threonine, 87-valine, 88-asparagine, 90-
     threonine] (human) (9CI)
                             (CA INDEX NAME)
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
RETABLE
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de Valk, B
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Ernst, S
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Silverberg, D
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L108 ANSWER 3 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN
AN
     2004:333839 HCAPLUS
DN
     140:352406
TI
    Erythropoietin glycosylation and the modification of
    protein structure and activity for therapeutic use
    De Frees, Shawn; Zopf, David; Bayer, Robert; Bowe, Caryn; Hakes, David;
IN
    Chen, Xi
    Neose Technologies, Inc., USA
PA
    PCT Int. Appl., 1018 pp.
SO
    CODEN: PIXXD2
DT
    Patent
LA
    English
FAN.CNT 15
    PATENT NO.
                       KIND
                              DATE
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    WO 2003031464
                       A2
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         RW: GH, GH, GM, GM, KE, KE, LS, LS, MW, MW, MZ, MZ, SD, SD, SL, SL,
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             RU, TJ, TM, AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB,
             GR, IE, IT, LU, MC, NL, PT, SE, SK, TR, BF, BJ, CF, CG, CI, CM,
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                                              US 2002-287994
     US 2004137557
                           A1
                                 20040715
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     CA 2501832
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PRAI WO 2002-US32263
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                                 20030219
     US 2003-410945
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     US 2001-328523P
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     US 2001-344692P
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     US 2002-387292P
                           p
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                           Ρ
                                 20020625
     US 2002-396594P
                           Ρ
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     US 2002-404249P
                           P
                                 20020816
     US 2002-407527P
                           Ρ
                                 20020828
     WO 2003-US31974
                           W
                                 20031008
     The invention includes methods and compns. for remodeling a peptide mol.,
\mathbf{AB}
     including the addition or deletion of one or more glycosyl groups to a
     peptide, and/or the addition of a modifying group to a peptide. Methods of
     modifying the structure and properties of erythropoietin by
     introduction of glycosidation are described. The method uses substitution
     variants of erythropoietin to introduce sites that can be
     glycosylated enzymically. The primary glycosylation may
     then be used to add further sugar residues. The glycosidation, which may include the introduction of N-acetylglucose, N-acetylgalactose, and sialic
     acid and mannosyl and fucosyl oligosaccharides. The carbohydrate moiety
     may in turn be modified by PEGylation. A biantennary
     glycosidated derivative of Epogen had 146% of the activity of the unmodified
     protein. The glycosylated proteins had longer serum half-lives
     than the unmodified protein and showed longer term effects on blood Hb
     levels.
IT
     681860-67-3DP, substitution derivs., glycosylated,
     PEGylated
     RL: BPN (Biosynthetic preparation); PRP (Properties); THU (Therapeutic
     use); BIOL (Biological study); PREP (Preparation); USES (Uses)
         (amino acid sequence; erythropoietin glycosylation
        and modification of protein structure and activity for therapeutic use)
RN
     681860-67-3 HCAPLUS
CN
     Erythropoietin (human 165-amino acid isoform) (9CI) (CA INDEX NAME)
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
IT
     11096-26-7DP, Erythropoietin, glycosylated
     derivs. 25322-68-3DP, Polyethylene glycol,
     reaction products with glycosylated erythropoietin
     RL: BPN (Biosynthetic preparation); PRP (Properties); THU (Therapeutic
     use); BIOL (Biological study); PREP (Preparation); USES (Uses)
        (erythropoietin glycosylation and modification of
        protein structure and activity for therapeutic use)
RN
     11096-26-7 HCAPLUS
```

(CA INDEX NAME)

CN

Erythropoietin (9CI)

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*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
       25322-68-3 HCAPLUS
       Poly(oxy-1,2-ethanediyl), \alpha-hydro-\omega-hydroxy- (9CI)
                                                                            (CA INDEX
       NAME)
но Сн2-Сн2-О н
 IT
       113427-24-0DP, Epogen, glycosylated derivs.
       RL: PKT (Pharmacokinetics); PNU (Preparation, unclassified); BIOL
       (Biological study); PREP (Preparation)
            (preparation and pharmacokinetics of; erythropoietin
           glycosylation and modification of protein structure and
           activity for therapeutic use)
       113427-24-0 HCAPLUS
RN
CN
       1-165-Erythropoietin (human clone \( \lambda \text{HEPOFL13} \) protein moiety),
       glycoform \alpha (9CI) (CA INDEX NAME)
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
L108 ANSWER 4 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN
       2004:287861 HCAPLUS
AN
DN
       140:320038
       Chimeric and humanized anti-granulocyte antibodies,
TT
       immunoconjugates and labeled antibodies for diagnosis and
       treatment of malignancy, infection and inflammation
IN
       Goldenberg, David M.; Hansen, Hans; Leung, Shui-on
PA
       Immunomedics, Inc., USA; Mccall, John Douglas
SO
       PCT Int. Appl., 134 pp.
       CODEN: PIXXD2
DT
       Patent
LΑ
       English
FAN.CNT 1
       PATENT NO.
                                 KIND
                                                          APPLICATION NO.
                                           DATE
                                                                                        DATE
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       WO 2004029093
                                  A2
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                                                          WO 2003-GB4229
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       WO 2004029093
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           2004029093

A3 20040603

W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NI, NO, NZ, OM, PG, PH, PL, PT, RO, RU, SC, SD, SE, SG, SK, SL, SY, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, YU, ZA, ZM, ZW

RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM, AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HU, IE, IT, LU, MC, NL, PT, RO, SE, SI, SK, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG
       CN 1542019
                                           20041103
                                  Α
                                                       CN 2003-123054
                                                                                         20030429 <--
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                                                        CA 2003 22
EP 2003-751001
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      EP 1546204
                                           20050629
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                 IE, SI, LT, LV, FI, RO, MK, CY, AL, TR, BG, CZ, EE, HU, SK
PRAI US 2002-414341P
                                  P
                                           20020930
                                                        <--
      WO 2003-GB4229
                                  W
                                           20030930
       The present invention provides humanized, chimeric and human MN3
AB
      antibodies, fusion proteins, and fragments that bind NCA90 and NCA95
       antigens. The antibodies, fusion proteins, and fragments thereof, as well
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as combinations with other suitable antibodies, are useful for the
     treatment and diagnosis of granulocyte related disorders and diseases,
     such as leukemia.
ΙT
     11096-26-7, Erythropoietin
     RL: BSU (Biological study, unclassified); DGN (Diagnostic use); PRP
     (Properties); THU (Therapeutic use); BIOL (Biological study); USES (Uses)
        (chimeric and humanized anti-granulocyte antibodies,
        immunoconjugates and labeled antibodies for diagnosis and
        treatment of malignancy, infection and inflammation)
RN
     11096-26-7 HCAPLUS
CN
     Erythropoietin (9CI) (CA INDEX NAME)
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
    7439-89-6, Iron, biological studies 15438-31-0
     , Iron(2+), biological studies 20074-52-6,
     Iron(3+), biological studies
     RL: BSU (Biological study, unclassified); DGN (Diagnostic use); THU
     (Therapeutic use); BIOL (Biological study); USES (Uses)
        (chimeric and humanized anti-granulocyte antibodies,
        immunoconjugates and labeled antibodies for diagnosis and
        treatment of malignancy, infection and inflammation)
RN
     7439-89-6 HCAPLUS
     Iron (7CI, 8CI, 9CI) (CA INDEX NAME)
Fe
RN
     15438-31-0 HCAPLUS
     Iron, ion (Fe2+) (8CI, 9CI) (CA INDEX NAME)
CN
Fe2+
     20074-52-6 HCAPLUS
RN
     Iron, ion (Fe3+) (8CI, 9CI) (CA INDEX NAME)
CN
Fe3+
L108 ANSWER 5 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN
AΝ
     2004:203692 HCAPLUS
DN
     140:229921
ΤI
    Use of erythropoietin and analogs to treat disturbances of iron
     distribution in diabetes
IN
     Lehmann, Paul; Roeddiger, Ralf; Walter-Matsui, Ruth
PA
     F. Hoffmann-La Roche A.-G., Switz.
SO
     PCT Int. Appl., 31 pp.
     CODEN: PIXXD2
DT
     Patent
LΑ
    English
FAN.CNT 1
     PATENT NO.
                        KIND
                                DATE
                                           APPLICATION NO.
                                                                   DATE
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                                           ------
PI
    WO 2004019972
                                20040311
                        A1
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        W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN,
             CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH,
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          PG, PH, PL, PT, RO, RU, SC, SD, SE, SG, SK, SL, SY, TJ, TM, TN, TR, TT, TZ, UA, UG, UZ, VC, VN, YU, ZA, ZM, ZW

RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM, AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HU, IE, IT, LU, MC, NL, PT, RO, SE, SI, SK, TR, RF RJ CF CG CJ CM GA GN GO GW ML MR NE SN TD TG
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      US 2004110679
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                                                                               20030820 <--
PRAI EP 2002-19100
                              Α
                                      20020829
                                                 <--
      WO 2003-EP9194
                                      20030820
      The present invention relates to the use of erythropoietin for
AB
      the treatment of disturbances of iron distribution in diabetes.
IT
      668496-68-2 668496-69-3
      RL: PAC (Pharmacological activity); PRP (Properties); THU (Therapeutic
      use); BIOL (Biological study); USES (Uses)
          (amino acid sequence; use of erythropoietin (Epo)
         and analogs to treat disturbances of iron distribution in diabetes)
RN
      668496-68-2 HCAPLUS
      Erythropoietin (human 165-amino acids variant) (9CI) (CA INDEX NAME)
CN
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
      668496-69-3 HCAPLUS
RN
      Erythropoietin (human 166-amino acids variant) (9CI) (CA INDEX NAME)
CN
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
IT
      11096-26-7, Erythropoietin 11096-26-7D,
      Erythropoietin, glycosylated and PEGylated
      variants and conjugates
      RL: PAC (Pharmacological activity); PRP (Properties); THU (Therapeutic
      use); BIOL (Biological study); USES (Uses)
          (use of erythropoietin (Epo) and analogs to treat
         disturbances of iron distribution in diabetes)
RN
      11096-26-7 HCAPLUS
CN
      Erythropoietin (9CI)
                                (CA INDEX NAME)
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
RN
      11096-26-7 HCAPLUS
CN
      Erythropoietin (9CI)
                                (CA INDEX NAME)
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
      113427-24-0, Epoetin alfa 122312-54-3
      , Epoetin beta 209810-58-2,
      Darbepoetin alfa
      RL: PAC (Pharmacological activity); THU (Therapeutic use); BIOL
      (Biological study); USES (Uses)
         (use of erythropoietin (Epo) and analogs to treat
         disturbances of iron distribution in diabetes)
RN
      113427-24-0 HCAPLUS
CN
      1-165-Erythropoietin (human clone λΗΕΡΟFL13 protein moiety),
      glycoform \alpha (9CI) (CA INDEX NAME)
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
RN
      122312-54-3 HCAPLUS
CN
      1-165-Erythropoietin (human clone λΗΕΡΟFL13 protein moiety),
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glycoform  $\beta$  (9CI) (CA INDEX NAME)

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

RN 209810-58-2 HCAPLUS

CN Erythropoietin [30-asparagine, 32-threonine, 87-valine, 88-asparagine, 90-threonine] (human) (9CI) (CA INDEX NAME)

## \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

#### RETABLE

Referenced Author (RAU)	Year (RPY)	 	Referenced Work (RWK)	Referenced File
Harold, T	2002	 	US 6440932 B1	HCAPLUS
Hoffmann La Roche	2001		WO 0187329 A	HCAPLUS
Hoffmann La Roche	2003		WO 03025583 A	HCAPLUS

L108 ANSWER 6 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN

AN 2003:282607 HCAPLUS

DN 138:298131

TI PEGylated and diglycosylated erythropoietin

with improved pharmaceutical properties in induction of erythropoiesis

IN Tischer, Wilhelm

PA F. Hoffmann-La Roche Ag, Switz.

SO PCT Int. Appl., 22 pp.

CODEN: PIXXD2

DT Patent

LA English

FAN.CNT 1

FAN.CNI I																			
						KIND DATE			APPLICATION NO.					DATE					
ΡI						_					WO 2	002-1	EPIO:	556		20	0020	920 <-	_
	WO	2003	0292:	91		<b>A</b> 3		2003	0724		•								
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			GM.	HR.	HU.	ID.	IL.	IN,	ıs.	JP.	KE.	KG.	KP.	KR.	KZ.	LC.	LK.	LR.	
			-					MD,		•	-					-	-	-	
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			•		-			TM,		•	•		•	•		•		•	
			FI,	FR,	GB,	GR,	ΙE,	IT,	LU,	MC,	NL,	PT,	SE,	SK,	TR,	BF,	ВJ,	CF,	
			CG,	CI,	CM,	GΑ,	GN,	GQ,	GW,	ML,	MR,	ΝE,	SN,	TD,	TG				
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	US	6930	086			B2		2005	0816										
	CA	2460	489			AA		2003	0410		CA 2	002-	2460	489		20	0020	920 <-	. <b>_</b>
																		920 <-	
																		PT,	
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	CINT	1550		•				RO,					•		•				
		1558																920 <-	
												003-	5325	36		20	0020	920 <-	-
PRAI	ΕP	2001	-122	555		Α		2001	0925	< -	-			-					
	WO	2002	-EP1	0556		W		2002	0920	<-	-								
								-								-			

AB The invention provides a new class of EPO muteins with improved pharmaceutical properties. The EPO muteins according to the invention have the in vivo biol. activity of causing bone marrow cells to increase production of reticulocytes and red blood cells. The invention provides an erythropoietin mutein which has retained the potential N-glycosylation sites at Asn24, Asn38, Asn83, is N-glycosylated at Asn38 and Asn83 but is not N-glycosylated at Asn24 and is preferably linked at the N-terminal amino group and/or the



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ε-amino group of Lys20 to poly(ethylene
     glycol) group(s) (PEG), preferably to
     alkoxypoly(ethylene glycol) group(s), more preferably to lower
     methoxypoly(ethylene glycol) group(s). The muteins of this invention have
     the same uses as EPO. In particular, the muteins of this
     invention are useful to treat patients by stimulating the division and
     differentiation of committed erythroid progenitors in the bone marrow.
     The present invention also includes a method for the treatment of anemia
     in humans and the use of the muteins for the manufacturing of a pharmaceutical
     agent preferably for such treatment. The present invention also includes
     a method for preparing erythropoietin muteins according to the
     invention, which comprises the production of a glycosylated
     EPO fragment consisting of the amino acids 26-165-(EPO
     26-165) and subsequent fusion of said fragment with a
     nonglycosylated but preferably PEGylated EPO
     fragment consisting of the amino acids 1-28 (EPO 1-28).
     510776-46-2DP, muteins 510776-47-3DP, muteins
IT
     RL: BPN (Biosynthetic preparation); PAC (Pharmacological activity); SPN
     (Synthetic preparation); THU (Therapeutic use); BIOL (Biological study);
     PREP (Preparation); USES (Uses)
        (amino acid sequence; preparation of PEGylated and
        diglycosylated erythropoietin with improved
        pharmaceutical properties in induction of erythropoiesis)
RN
     510776-46-2 HCAPLUS
CN
     Erythropoietin (human 165-amino acid isoform) (9CI)
                                                          (CA INDEX NAME)
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
RN
     510776-47-3 HCAPLUS
CN
     Erythropoietin (human 166-amino acid isoform) (9CI)
                                                          (CA INDEX NAME)
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
IT
     510776-48-4, 29-165-erythropoietin (human)
     RL: RCT (Reactant); RACT (Reactant or reagent)
        (amino acid sequence; preparation of PEGylated and
        diglycosylated erythropoietin with improved
        pharmaceutical properties in induction of erythropoiesis)
RN
     510776-48-4 HCAPLUS
CN
     29-165-erythropoietin (human) (9CI) (CA INDEX NAME)
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
IT
     11096-26-7DP, Erythropoietin, muteins
     RL: BPN (Biosynthetic preparation); PAC (Pharmacological activity); SPN
     (Synthetic preparation); THU (Therapeutic use); BIOL (Biological study);
     PREP (Preparation); USES (Uses)
        (preparation of PEGylated and diglycosylated
        erythropoietin with improved pharmaceutical properties in
        induction of erythropoiesis)
RN
     11096-26-7 HCAPLUS
     Erythropoietin (9CI) (CA INDEX NAME)
CN
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
L108 ANSWER 7 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN
AN
     2002:869575 HCAPLUS
DN
     137:346941
TI
     Method for improving the quality of life of patients by administration of
     erythropoietin (RhuEPO)
IN
     Zaharia, Veronica C.
PΑ
SO
     U.S. Pat. Appl. Publ., 4 pp., Cont.-in-part of U.S. Ser. No. 872,630.
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CODEN: USXXCO DT Patent LΑ English FAN.CNT 3 PATENT NO. KIND DATE APPLICATION NO. DATE ----\_\_\_\_\_ -----\_\_\_\_\_ US 2002169129 20021114 PΙ **A**1 US 2002-133545 20020426 <--19990914 US 5951996 Α US 1998-18815 19980204 <--US 6274158 В1 20010814 US 1999-335076 19990617 <--US 6521245 B1 20030218 US 2001-872630 20010601 <--PRAI US 1998-18815 A2 19980204 <--US 1998-91598P Р 19980702 <--Ρ US 1999-125253P 19990319 <--US 1999-335076 **A3** 19990617 <--US 2001-287206P P 20010428 <--US 2001-872630 A2 20010601 <--AB A method for providing various benefits with the administration of different quantities of Erythropoietin. The method provides for enhancing the of quality of life by administration of Erythropoietin before a substantial increases in Hb occurs. improvement in quality of life is independent of the hemopoietic effect. In larger quantities the administration of RhuEPO leads to repair of vascular damage and leads to the redistribution of the iron trapped in storage organs, from where it cannot be used for red blood cell production, into the hemopoietic system leading to enhanced red blood cell production TT 11096-26-7, Erythropoietin RL: PAC (Pharmacological activity); THU (Therapeutic use); BIOL (Biological study); USES (Uses) (method for improving the quality of life of patients by administration of erythropoietin (RhuEPO)) RN11096-26-7 HCAPLUS Erythropoietin (9CI) (CA INDEX NAME) CN \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\* 7439-89-6, Iron, biological studies RL: BSU (Biological study, unclassified); BIOL (Biological study) (method for limiting chronic blood loss by administering RhuEPO to prevent iron loss and to increase Hb level, increased mean corpuscular Hb, and increased red blood cell hemoglobinization.) 7439-89-6 HCAPLUS RNIron (7CI, 8CI, 9CI) CN (CA INDEX NAME)

Fe

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L108 ANSWER 8 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN
AN
     2002:785122 HCAPLUS
DN
     138:298038
     Long-term reversal of chronic anemia using a hypoxia-regulated
ΤI
     erythropoietin gene therapy
     Binley, Katie; Askham, Zoe; Iqball, Sharifah; Spearman, Hayley; Martin,
AU
     Leigh; de Alwis, Mahesh; Thrasher, Adrian J.; Ali, Robin R.; Maxwell,
     Patrick H.; Kingsman, Susan; Naylor, Stuart
     Oxford BioMedica (UK) Ltd, London, OX4 4GA, UK
CS
     Blood (2002), 100(7), 2406-2413
SO
     CODEN: BLOOAW; ISSN: 0006-4971
     American Society of Hematology
PΒ
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DT Journal

LΑ English

AB Anemia is a common clin. problem, and there is much interest in its role in promoting left ventricular hypertrophy through increasing cardiac workload. Normally, red blood cell production is adjusted through the regulation of erythropoietin (Epo) production by the kidney. One important cause of anemia is relative deficiency of Epo, which occurs in most types of renal disease. Clin., this can be corrected by supplementation with recombinant Epo. Here the authors describe an oxygen-regulated gene therapy approach to treating homozygous erythropoietin-SV40 T antigen (Epo-TAgh) mice with relative erythropoietin deficiency. The authors used vectors in which murine Epo expression was directed by an Oxford Biomedica hypoxia response element (OBHRE) or a constitutive cytomegalovirus (CMV) promoter. Both corrected anemia, but CMV-Epo -treated mice acquired fatal polycythemia. In contrast, OBHRE-Epo corrected the hematocrit level in anemic mice to a normal physiol. level that stabilized without resulting in polycythemia. Importantly, the OBHRE-Epo vector had no significant effect on the hematocrit of control mice. Homozygous Epo-TAgh mice display cardiac hypertrophy, a common adaptive response in patients with chronic anemia. In the OBHRE-Epo-treated Epo-TAgh mice, the authors observed a significant reversal of cardiac hypertrophy. The authors conclude that the OBHRE promoter gives rise to physiol. regulated Epo secretion such that the hematocrit level is corrected to healthy in anemic Epo-TAgh mice. This establishes that a hypoxia regulatory mechanism similar to the natural mechanism can be achieved, and it makes EPO gene therapy more attractive and safer in clin. settings. The authors envisage that this control system will allow regulated delivery of therapeutic gene products in other ischemic settings. IT 7439-89-6, Iron, biological studies

RL: BSU (Biological study, unclassified); BIOL (Biological study) (long-term reversal of chronic anemia using hypoxia-regulated erythropoietin gene therapy)

RN

7439-89-6 HCAPLUS Iron (7CI, 8CI, 9CI) CN (CA INDEX NAME)

Fe ·

#### IT 11096-26-7, Erythropoietin

RL: BSU (Biological study, unclassified); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

(long-term reversal of chronic anemia using hypoxia-regulated erythropoietin gene therapy)

RN 11096-26-7 HCAPLUS

Erythropoietin (9CI) (CA INDEX NAME)

### \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

RETABLE

Referenced Author (RAU)	,	VOL (RVL)		Referenced Work (RWK)	Referenced File
Anon Bachmann, S Bartholomew, A Beall, C Binley, K Boast, K	1989 1993	2 41	20	Lancet J Histochem Cytochem Hum Gene Ther Gene Ther Gene Ther Hum Gene Ther	

n-1-1 n	1	1.00		1	
Bohl, D	1998	92	1512	Blood	HCAPLUS
Bohl, D	2000	95	2793	Blood	HCAPLUS
Bohl, D	1997	3	299	Nat Med	HCAPLUS
Bron, D	2001	28	1	Semin Oncol	MEDLINE
Cowgill, L	1998	212	521	J Am Vet Med Assoc	HCAPLUS
Dalle, B	1999	6	157	Gene Ther	HCAPLUS
Erslev, A	1985	41	213	Nephron	MEDLINE
Eschbach, J	1989	35	134	Kidney Int	MEDLINE
Foley, R	1995	5	2024	J Am Soc Nephrol	MEDLINE
Goodnough, L	2000	96	823	Blood	HCAPLUS
Griffiths, L	2000	7	255	Gene Ther	HCAPLUS
Hamamori, Y	1995	95	1808	J Clin Invest	HCAPLUS
Jelkmann, W	1992	72	449	Physiol Rev	HCAPLUS
Kina, T	2000	109	280	Br J Haematol	HCAPLUS
Krystal, G	1983	11	649	Exp Hematol	HCAPLUS
Lynch, C	1999	1	493	Curr Opin Mol Ther	HCAPLUS
Maxwell, P	1993	44	1149	Kidney Int	HCAPLUS
Maxwell, P	1993	90	2423	Proc Natl Acad Sci U	
Middleton, R	2001	12	1079	J Am Soc Nephrol	MEDLINE
Post, D		8	1801	Gene Ther	HCAPLUS
Raja, K	1997	96	248	Br J Haematol	HCAPLUS
Rendahl, K	1998	16	757	Nat Biotechnol	HCAPLUS
Rinsch, C	1997	8	1881	Hum Gene Ther	HCAPLUS
Rudich, S	2000	90	102	J Surg Res	HCAPLUS
Semenza, G	2000	14	1983	Genes Dev	HCAPLUS
Seppen, J	2001	98	594	Blood	HCAPLUS
Serquera, C	1999	10	375	Hum Gene Ther	HCAPLUS
Setoquch, Y	1994	84	2946	Blood	ICAPIOS
Villeval, J	1994	84	928	Blood	HCAPLUS
Wang, G	1993	90	4304	Proc Natl Acad Sci U	
Wang, G	1995	92	5510	Proc Natl Acad Sci U	
Ye, X	1999	283	88	Science	
Zhang, X	1999	10	2527	Hum Gene Ther	HCAPLUS
Zhou, S	1998	5	665	Gene Ther	HCAPLUS
2	1 1 9 9 0	1 2	1003	Gene Inet	HCAPLUS

L108 ANSWER 9 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN

AN 2002:761224 HCAPLUS

DN 137:288375

- TI The correction of anemia in severe resistant heart failure with erythropoietin and intravenous iron prevents the progression of both the heart and the renal failure and markedly reduces hospitalization
- AU Silverberg, D. S.; Wexler, D.; Blum, M.; Tchebiner, J.; Sheps, D.; Keren, G.; Schwartz, D.; Baruch, R.; Yachnin, T.; Shaked, M.; Zubkov, A.; Steinbruch, S.; Iaina, A.
- CS Department of Nephrology and Cardiology and Congestive Heart Failure Unit and Medical Department B, Tel Aviv Medical Center, Tel Aviv-Jaffa, 64239, Israel
- SO Clinical Nephrology (2002), 58(1, Suppl. 1), S37-S45 CODEN: CLNHBI; ISSN: 0301-0430
- PB Dustri-Verlag Dr. Karl Feistle
- DT Journal; General Review
- LA English
- AB A review. Both Congestive Heart Failure (CHF) and Chronic Renal Failure (CRF) are increasing steadily in the community. We propose that there is a vicious circle established whereby CHF and CRF both cause anemia and the anemia then worsens both the CHF and CRF causing more anemia and so on. We call this the Cardio Renal Anemia (CRA) syndrome. By the combination of active treatment of the CHF and control of the anemia with s.c. erythropoietin and i.v. iron, the progression of both



the CHF and the CRF can be slowed or stopped in most cases, the quality of life improved and the need for recurrent hospitalization reduced. This will involve cooperation between internists, cardiologists, and nephrologists to allow early and maximal therapy of both the CHF and the anemia.

TT 7439-89-6, Iron, biological studies 11096-26-7
, Erythropoietin

RL: PAC (Pharmacological activity); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

(erythropoietin and i.v. iron correction of anemia

in severe resistant heart failure patients prevents progression of both heart and renal failure and markedly reduces hospitalization)

RN 7439-89-6 HCAPLUS

CN Iron (7CI, 8CI, 9CI) (CA INDEX NAME)

Fe

RN 11096-26-7 HCAPLUS

CN Erythropoietin (9CI) (CA INDEX NAME)

# \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\* RETABLE

Referenced Author (RAU)	Year (RPY)	VOL	PG (RPG)	Referenced Work (RWK)	Referenced File
•	(RFI)    -====+		(RPG)  ======	! (KWK) +====================================	
Al-Ahmad, A	2001	38	955	J Am Coll Cardiol	MEDLINE
Anand, I	1993	70	357	Br Heart J	MEDLINE
Bosman, D	2001	24	495	Diabetes Care	MEDLINE
Capes, S	2000	23	377	Diabetes Care	HCAPLUS
Carson, J	1995	170	32S	Amer J Surg	
Cowie, M	1997	18	208	Eur Heart J	HCAPLUS
Dries, D	2001	38	421	J Am Coll Cardiology	MEDLINE
Felker, G	2001	17		Circulation	
Fine, L	1998	53	s74	Kidney Int	
Fine, L	2000	57	S22	Kidney Int	
Fishbane, S	1995	26	41	Am J Kidney Dis	MEDLINE
Foley, R	1996	28	53	Am J Kidney Dis	MEDLINE
Foley, R	1995	47	186	Kidney Int	MEDLINE
Goodnough, L	2001	345	1272	N Engl J Med	MEDLINE
Hampl, H	1985	39	102	Nephron	MEDLINE
He, J	2001	161	996	Arch Int Med	MEDLINE
Herrera-Garza, E	1999	115	1170	Chest	MEDLINE
Hillege, H	1999	10	A384	J Am Soc Nephrol	
Holland, D	2000	15	650	Nephrol Dial Transpl	MEDLINE
Iverson, P	2002	282	R166	Am J Physiol Regul I	
Jafar, T	2001	60	1131	Kidney Int	HCAPLUS
Johnson, D	1996	5	186	Current Opinion in N	MEDLINE
Jungers, P	2001	16	307	Nephrol Dialysis Tra	HCAPLUS
Katz, A	1994	121	363	Ann Intern Med	HCAPLUS
Knight, E	1999	138	849	Am Ht J	MEDLINE
Kratz, A	1998	339	1063	N Engl J Med	MEDLINE
Kuriyama, S	1997	77	176	Nephron	HCAPLUS
Linde, T	1996	30	115	Scand J Urol Nephrol	MEDLINE
Longenecker, J	2000	11	520	J Am Soc Nephrol	MEDLINE
Macdougall, I	1996	50	1694	Kidney Int	HCAPLUS
Macdougall, I	1999	14	1836	Nephrol Dial Transpl	MEDLINE
Major, A	1997	98	292	Brit J Haematol	HCAPLUS
Mancini, D	2000	101	1080	Circulation	MEDLINE



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Mancini, D
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McAlister, F
                         1999
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                                      87
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                                                                    MEDLINE
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                         1999
Means, R
                               70
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Miller, L
                         2001
                               19
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                                                                    MEDLINE
                                             Cardiology Clinics
Mosterd, A
                                      1318
                                             Eur Heart J
                         2001
                               22
                                                                    MEDLINE
O'Connell, J
                         2000
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Packer, M
                         1999
                                             Am J Cardiol
                               83
                                      1
Perry, H
                         1995
                               25
                                      587
                                             Hypertension
Petrie, M
                         2001
                               22
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                                             Eur Heart J
                                                                    MEDLINE
Philbin, E
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                                      605
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                                                                    MEDLINE
Silverberg, D
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                               35
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                                             J Am Coll Cardiol
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                         2000
                               53
                                      S23
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Sommerburg, O
Sunder-Plassmann, G
                         2001
                               38
                                      S20
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Tong, E
                         2001
                               21
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                                             Seminars in Nephrolo HCAPLUS
Tsuyuki, R
                         2001
                               161
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                                             Archives Int Med
                                                                    MEDLINE
US Renal Data System
                         2001
                                             Annual Data Report
                               38
Vaziri, N
                         2001
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                                                                    HCAPLUS
Wahr, J
                         1998
                                81
                                      10
                                             Br J Anaest
Winkler, A
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                         1999
                                16
                                             Diabetes Medicine
                                                                    MEDLINE
Wisniacki, N
                         2001
                               85
                                      P4
                                             Heart
Wu, W
                         2001
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                                                                    MEDLINE
Yoshida, H
                         1998
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L108 ANSWER 10 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN

AN 2002:487418 HCAPLUS

DN 137:68127

TI Erythropoietin conjugates

IN Burg, Josef; Engel, Alfred; Franze, Reinhard; Hilger, Bernd; Schurig, Hartmut Ernst; Tischer, Wilhelm; Wozny, Manfred

PA F. Hoffmann-La Roche Ag, Switz.

SO PCT Int. Appl., 40 pp.

CODEN: PIXXD2

DT Patent

LA English

FAN. CNT 1

	PATENT NO.		APPLICATION NO.	DATE		
PI	WO 2002049673 WO 2002049673	A2 20020627	WO 2001-EP14434	20011208 <		
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	EP 1345628	A2 20030924	EP 2001-984811	20011208 <		
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Page 20

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20040225
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                                             JP 2002-551010
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     US 2002115833
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                                             US 2001-14363
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     ZA 2003004647
                          Α
                                 20040913
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                                                                     20030613 <--
PRAI EP 2000-127891
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                          Α
                                           <--
     WO 2001-EP14434
                          W
                                 20011208
     The present invention refers to conjugates of
     erythropoietin with poly(ethylene
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AB

glycol) comprising an erythropoietin glycoprotein having an N-terminal  $\alpha$ -amino group and having the in vivo biol. activity of causing bone marrow cells to increase production of reticulocytes and red blood cells and selected from the group consisting of human erythropoietin and analogs thereof which have the sequence of human erythropoietin modified by the addition of from 1 to 6 glycosylation sites or a rearrangement of at least one glycosylation site; said glycoprotein being covalently linked to one poly(ethylene glycol) group of the formula -CO-(CH2)x-(OCH2CH2)m-OR with the -CO of the poly( ethylene glycol) group forming an amide bond with said N-terminal  $\alpha$ -amino group; wherein R is lower alkyl; x is 2 or 3; and m is from about 450 to about 1350.

IT 11096-26-7DP, Erythropoietin, conjugates

RL: PNU (Preparation, unclassified); THU (Therapeutic use); BIOL (Biological study); PREP (Preparation); USES (Uses)

(glycosylation site-augmented human erythropoietin conjugates with PEG)

11096-26-7 HCAPLUS RN

Erythropoietin (9CI) (CA INDEX NAME) CN

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

11096-26-7, Erythropoietin 25322-68-3D, Polyethylene glycol, erythropoietin conjugates

> RL: RCT (Reactant); RACT (Reactant or reagent) (glycosylation site-augmented human erythropoietin conjugates with PEG)

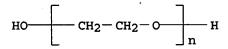
RN 11096-26-7 HCAPLUS

CN Erythropoietin (9CI) (CA INDEX NAME)

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

25322-68-3 HCAPLUS RN

Poly(oxy-1,2-ethanediyl), α-hydro-ω-hydroxy- (9CI) (CA INDEX CN



L108 ANSWER 11 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN

2002:409245 HCAPLUS AN

DN 136:380106

Method of treating congestive heart failure with erythropoietin TI and an iron compound

Iaina, Adrian; Wexler, Dov; Silverberg, Donald S. IN

PA Israel

U.S. Pat. Appl. Publ., 3 pp. SO

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CODEN: USXXCO
DT
      Patent
      English
LA
FAN.CNT 1
      PATENT NO.
                              KIND
                                      DATE
                                                   APPLICATION NO.
                                                                                 DATE
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                                                  US 2000-725161
PΤ
      US 2002065214
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                                      20020530
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PRAI US 2000-725161
                                      20001129 <--
AΒ
      A method of treating congestive heart failure in a subject suffering
      therefrom, comprising administering erythropoietin and i.v.
      administering an i.v. administrable iron compound to the subject.
      The iron is preferably administered in the form of a complex of
      a ferric hydroxide with erythropoietin.
IT
      11096-26-7, Erythropoietin
      RL: PAC (Pharmacological activity); THU (Therapeutic use); BIOL
      (Biological study); USES (Uses)
          (method of treating congestive heart failure with
         erythropoietin and an iron compound)
RN
      11096-26-7 HCAPLUS
CN
      Erythropoietin (9CI)
                                 (CA INDEX NAME)
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
L108 ANSWER 12 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN
      2001:850963 HCAPLUS
AN -
DN
      136:11065
ΤI
      New pharmaceutical composition
IN
      Papadimitriou, Apollon
      F. Hoffmann-La Roche A.-G., Switz.
PA
SO
      PCT Int. Appl., 64 pp.
      CODEN: PIXXD2
DT
      Patent
      English
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FAN.CNT 1
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      PATENT NO.
                              KIND
                                      DATE
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      EP 1525889
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US 2001-853731

ZA 2002-8500

20010511 <--

20021021 <--

20020328

20040128

US 2002037841

ZA 2002008500

**A1** 

Α

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

25322-68-3D, Polyethylene glycol, protein IT conjugates

RL: PEP (Physical, engineering or chemical process); THU (Therapeutic use); BIOL (Biological study); PROC (Process); USES (Uses) (stabilized erythropoietin pharmaceutical composition)

RN 25322-68-3 HCAPLUS

CN Poly(oxy-1,2-ethanediyl), α-hydro-ω-hydroxy- (9CI) (CA INDEX

HO 
$$CH_2-CH_2-O$$
 H

### RETABLE

AB

for

IT

RN

CN

RN

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IT

RN

CN

Referenced Author |Year | VOL | PG Referenced Work Referenced (RAU) | (RPY) | (RVL) | (RPG) | File (RWK)



L108 ANSWER 13 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN

AN 2001:715798 HCAPLUS

DN 136:15603

TI Erythropoietin therapy and preoperative autologous blood donation in children undergoing open heart surgery

- AU Sonzogni, V.; Crupi, G.; Poma, R.; Annechino, F.; Ferri, F.; Filisetti, P.; Bellavita, P.
- CS Department of Anesthesiology, Ospedali Riuniti di Bergamo, Bergamo, Italy
- SO British Journal of Anaesthesia (2001), 87(3), 429-434 CODEN: BJANAD; ISSN: 0007-0912
- PB Oxford University Press
- DT Journal
- LA English
- AB We assessed the feasibility and efficacy of s.c. erythropoietin alpha (EPO) therapy and preoperative autologous blood donation (ABD) in children undergoing open heart surgery. Thirty-nine children were treated consecutively with EPO (100 U kg-1 s.c. three times a week in the 3 wk preceding the operation and i.v. on the day of surgery) and two ABDs were made (Group 1). As controls to compare transfusion requirements, 39 consecutive age-matched patients who had undergone open heart surgery during the two preceding years were selected (Group 2). In a mean time of 20 (SD 5) days, 96% of scheduled ABDs were performed and only three mild vasovagal reactions were observed The mean volume of autologous red blood cells (RBC) collected was 6 (1) ml kg-1 and the mean volume of autologous RBC produced as a result of EPO therapy before surgery was 7 (3) ml kg-1, corresponding to a 28 (11)% increase in circulating RBC volume The mean volume of autologous RBC collected was not different from that produced [6 (1) vs. 7 (3) ml kg-1, P=0.4]. Allogenic blood was administered to three out of 39 children in Group 1 (7.7%) and to 24 out of 39 (61.5%) in Group 2. Treatment with s.c. EPO increases the amount of autologous blood that can be collected and minimizes allogenic blood exposure in children undergoing open heart surgery.
- IT 113427-24-0, Eprex

RL: PAC (Pharmacological activity); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

(erythropoietin therapy and preoperative autologous blood donation in children undergoing open heart surgery)

RN 113427-24-0 HCAPLUS

CN 1-165-Erythropoietin (human clone  $\lambda$ HEPOFL13 protein moiety), glycoform  $\alpha$  (9CI) (CA INDEX NAME)

# \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\* RETABLE

Referenced Author (RAU)	,	VOL (RVL)		Referenced Work   (RWK)	Referenced File
Adamson, J	1994	115	7	Surgery	MEDLINE
Anon	1988	260	2700	Consensus conference	
Baron, J	1997	33	64	Semin Hematol	
Beguin, Y	1999	84	541	Haematologica	HCAPLUS
Chaplin, H	1953	32	1309	J Clin Invest	
Cooley, D	1995	170	53	Am J Surg	
Coyle, D	2000	18	161	Pharmacoeconomics	MEDLINE

Despotis, G         1999         11         84         Semin Thorac Cardiov MEDLINE         MEDLINE           Fukahara, K         1994         114         504         J Thoroc Cardiovasc         MEDLINE           Goodnough, L         1994         101         354         Am J Clin Pathol         MEDLINE           Goodnough, L         1995         60         473         Ann Thorac Surg         MEDLINE           Goodnough, L         1999         321         1163         N Engl J Med         MEDLINE           Goodnough, L         1997         336         933         N Engl J Med         MEDLINE           Goodnough, L         1999         340         438         N Engl J Med         MEDLINE           Goodnough, L         1993         33         944         Transfusion         MEDLINE           Goodnough, L         1993         33         944         Transfusion         MEDLINE           Goodnough, L         1993         34         Transfusion         MEDLINE           Goodnough, L         1993         34         Transfusion         MEDLINE           Goodnough, L         1993         34         Transfusion         MEDLINE           Klaapper, E         1995         110				_		
Goodnough, L Goodn	<u> </u>		,	!		MEDLINE
Goodnough, L Goodn	- · · · · · · · · · · · · · · · · · · ·	1997	114	504	J Thoroc Cardiovasc	MEDLINE
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Goodnough, L         1989         321         1163         N Engl J Med         MEDLINE           Goodnough, L         1997         336         933         N Engl J Med         MEDLINE           Goodnough, L         1999         340         438         N Engl J Med         MEDLINE           Goodnough, L         1992         32         441         Transfusion         MEDLINE           Goodnough, L         1993         33         944         Transfusion         MEDLINE           Goodnough, L         1994         34         66         Transfusion         MEDLINE           Goodnough, L         1996         62         1955         Ann Thorac Surg         MEDLINE           Klapper, E         1995         110         1594         J Thorac Cardiovasc         MEDLINE           Krantz, S         1991         77         419         Blood         HCAPLUS           Marchetti, M	Goodnough, L	1995	60	473	Ann Thorac Surg	MEDLINE
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DN 135:267638

- TI The effect of correction of mild anemia in severe, resistant congestive heart failure using subcutaneous **erythropoietin** and intravenous **iron**: a randomized controlled study
- AU Silverberg, Donald S.; Wexler, Dov; Sheps, David; Blum, Miriam; Keren, Gad; Baruch, Ron; Schwartz, Doron; Yachnin, Tatyana; Steinbruch, Shoshana; Shapira, Itzhak; Laniado, Shlomo; Iaina, Adrian
- CS Department of Nephrology and Cardiology and Congestive Heart Failure, Tel Aviv Medical Center, Tel Aviv-Jaffa, Israel
- SO Journal of the American College of Cardiology (2001), 37(7), 1775-1780
  CODEN: JACCDI; ISSN: 0735-1097
- PB Elsevier Science Inc.
- DT Journal
- LA English
- This is a randomized controlled study of anemic patients with severe congestive heart failure (CHF) to assess the effect of correction of the anemia on cardiac and renal function and hospitalization. Although mild anemia occurs frequently in patients with CHF, there is very little information about the effect of correcting it with erythropoietin (EPO) and i.v. iron. Thirty-two patients with moderate to severe CHF (New York Heart Association [NYHA] class III to IV) who had a left ventricular ejection fraction (LVEF) of ≤40% despite maximally tolerated doses of CHF medications and whose Hb levels were persistently between 10.0 and 11.5 g% were randomized into two groups. Group A (16 patients) received s.c. EPO and IV iron to

increase the level of Hb to at least 12.5 g%. In Group B (16 patients) the anemia was not treated. The doses of all the CHF medications were maintained at the maximally tolerated levels except for oral and i.v. (IV) furosemide, whose doses were increased or decreased according to the clin. need. Over a mean of 8.2±2.6 mo, four patients in Group B and none in Group A died of CHF-related illnesses. The mean NYHA class improved by 42.1% in A and worsened by 11.4% in B. The LVEF increased by 5.5% in A and decreased by 5.4% in B. The serum creatinine did not change in A and increased by 28.6% in B. The need for oral and IV furosemide decreased by 51.3% and 91.3% resp. in A and increased by 28.5% and 28.0% resp. in B. The number of days spent in hospital compared with the same period of time before entering the study decreased by 79.0% in A and increased by 57.6% in B. When anemia in CHF is treated with EPO and IV iron, a marked improvement in cardiac and patient function is seen, associated with less hospitalization and renal impairment and less need for diuretics.

#### IT 11096-26-7, Erythropoietin

RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

(effect of correction of mild anemia in congestive heart failure using s.c. erythropoietin and i.v. iron)

RN 11096-26-7 HCAPLUS

CN Erythropoietin (9CI) (CA INDEX NAME)

# \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\* RETABLE

Referenced Author (RAU)	Year (RPY)	VOL (RVL)	PG (RPG)	Referenced Work (RWK)	Referenced
Al-Ahmad, A	2000	11	137A	J Am Soc Nephrol	
Alexander, M	1999	137	919	Am Heart J	MEDLINE
Ali, A	1999	138	1133	Am Heart J	MEDLINE
Anand, I	1993	70	357	Br Heart J	MEDLINE
Anon	1997	30	S193	Am J Kidney Dis	HIDDEINE
Bardaji, A	1998	32	970	Am J Kid Dis	MEDLINE
Carson, J	1995	170	32	Am J Surg	
Carson, J	1996	348	1055	Lancet	MEDLINE
De Simone, G	2000	101	152	Circulation	MEDLINE
Fine, L	1998	53	S74	Kidney Intl	
Fishbane, S	1995	26	41	Am J Kidney Dis	MEDLINE
Foley, R	1996	28	53	Am J Kidney Dis	MEDLINE
Ghali, J	1988	148	2013	Arch Intern Med	MEDLINE
Goldberg, N	1992	124	424	Am Heart J	MEDLINE
Haber, H	1991	324	353	N Engl J Med	MEDLINE
Harnett, J	1995	47	884	Kidney Int	MEDLINE
Hebert, P	1997	155	1618	Am J Respir Crit Car	
Hillege, H	1999	10	A384	J Am Soc Nephrol	
Jensen, J	1993	233	125	J Int Med	MEDLINE
Kannel, W	1987	8	23	Eur Heart J	
Knight, E	1999	138	849	Am Heart J	MEDLINE
Kuriyama, S	1996	9	426	Am J Hypertension	HCAPLUS
Locatelli, F	1998	13	1642	Nephrol Dial Transpl	MEDLINE
Low-Friedrich, I	1991	11	54	Am J Nephrol	MEDLINE
Ma, J	1999	10	610	J Am Soc Nephrol	MEDLINE
Macdougall, I	1996	50	1694	Kidney Int	HCAPLUS
Magri, P	1998	98	2849	Circulation	HCAPLUS
Maschio, G	1995	10	74	Nephrol Dial Transpl	
Opasich, C	1996	78	354	Am J Cardiol	MEDLINE
Packer, M	1999	83	1	Am J Cardiol	

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Perry, H
                       1995
                              25
                                    587
                                           Hypertension
Rich, M
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                                           J Am Geriatr Soc
                              44
                                     638
                                                                 MEDLINE
Silverberg, D
                        2000
                                    1737
                                           J Am Coll Cardiol
                              35
                                                                 HCAPLUS
Silverberg, D
                        1999
                              55
                                    S79
                                           Kidney Int
Silverberg, D
                        1996
                              72
                                    413
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                        1997
                              129
                                    97
                                            J Lab Clin Med
                                                                 HCAPLUS
Stefanski, A
                        1996
                              50
                                    1321
                                           Kidney Int
                                                                 MEDLINE
Wackers, F
                        1979
                              43
                                           Am J Cardiol
                                    1159
                                                                 MEDLINE
Wald, M
                        1996
                              167
                                    461
                                            J Cell Physiol
                                                                 HCAPLUS
Wald, M
                        1995
                              71
                                    190
                                           Nephron
                                                                 HCAPLUS
Wu, H
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                              126
                                    3597
                                           Development
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Xia, H
                                            J Am Soc Nephrol
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                              10
                                    1309
                                                                 MEDLINE
Yoshida, H
                        1998
                              153
                                    880
                                           Kidney Int
                                                                 MEDLINE
L108 ANSWER 15 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN
     2001:31360 HCAPLUS
DN
     134:105827
TI
     Erythropoietin derivatives
IN
     Burg, Josef; Hilger, Bernd; Josel, Hans-Peter
     F. Hoffmann-La Roche A.-G., Switz.
PA
     PCT Int. Appl., 40 pp.
SO
     CODEN: PIXXD2
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     English
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     PATENT NO.
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     EP 1196443
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AB
     Erythropoietin glycoprotein conjugates are disclosed,
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said conjugates comprise an erythropoietin



glycoprotein having at least one free amino group and having the in vivo biol. activity of causing bone marrow cells to increase production of reticulocytes and red blood cells and selected from the group consisting of human erythropoietin and analogs thereof which have the primary structure of human erythropoietin modified by the addition of from 1 to 6 glycosylation sites or by the rearrangement of at least one glycosylation site; said glycoprotein being covalently linked to form one to three lower-alkoxy poly(ethylene glycol) groups, each poly(ethylene glycol) group being covalently linked to the glycoprotein via a linker of the formula -C(0)-X-S-Y- with the C(0) of the linker forming an amide bond with one of said amino groups, wherein X and Y are as defined in the description and claims, the average mol. weight of each poly( ethylene glycol) moiety is from about 20 kilodaltons to about 40 kilodaltons, and the mol. weight of the conjugate is from about 51 kilodaltons to about 175 kilodaltons. 96024-34-9, Erythropoietin (human clone λΗΕΡΟFL13 protein moiety reduced) 134547-95-8, 1-165-Erythropoietin (human clone AHEPOFL13 protein moiety reduced) RL: BOC (Biological occurrence); BSU (Biological study, unclassified); PEP (Physical, engineering or chemical process); PRP (Properties); THU (Therapeutic use); BIOL (Biological study); OCCU (Occurrence); PROC (Process); USES (Uses) (amino acid sequence; erythropoietin derivs. for increasing production of erythrocytes and reticulocytes) RN 96024-34-9 HCAPLUS CN Erythropoietin (human clone λΗΕΡΟFL13 protein moiety reduced) (9CI) (CA INDEX NAME) \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\* RN134547-95-8 HCAPLUS CN 1-165-Erythropoietin (human clone \( \lambda \text{HEPOFL13} \) protein moiety reduced) (9CI) (CA INDEX NAME) \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\* IT 11096-26-7D, Erythropoietin, conjugates 25322-68-3D, erythropoietin conjugates RL: PEP (Physical, engineering or chemical process); THU (Therapeutic use); BIOL (Biological study); PROC (Process); USES (Uses) (erythropoietin derivs. for increasing production of erythrocytes and reticulocytes) RN 11096-26-7 HCAPLUS CN Erythropoietin (9CI) (CA INDEX NAME) \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\* RN25322-68-3 HCAPLUS CN Poly(oxy-1,2-ethanediyl),  $\alpha$ -hydro- $\omega$ -hydroxy- (9CI) (CA INDEX

$$HO = \begin{bmatrix} CH_2 - CH_2 - O \end{bmatrix}_n H$$

L108 ANSWER 16 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN ΑN 2001:10610 HCAPLUS DN 134:91083

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Erythropoietin derivatives for increasing bone marrow production
ΤI
     of reticulocytes and erythrocytes
IN
     Bailon, Pascal Sebastian
PA
     F. Hoffmann-La Roche A.-G., Switz.
SO
     Eur. Pat. Appl., 16 pp.
     CODEN: EPXXDW
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     English
LA
FAN.CNT 5
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                                                                   DATE
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AB
     The present invention refers to conjugates of
     erythropoietin with poly(ethylene
     glycol) comprising an erythropoietin glycoprotein having
     at least one free amino group and having the in vivo biol. activity of
     causing bone marrow cells to increase production of reticulocytes and red
     blood cells and selected from the group consisting of human
     erythropoietin and analogs thereof which have sequence of human
     erythropoietin modified by the addition of 1-6 glycosylation
     sites or a rearrangement of at least one glycosylation site;
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said glycoprotein being covalently linked to "n" poly(

-CO-(CH2)x(OCH2CH2)m-OR with the carbonyl of each poly(

ethylene glycol) groups of the formula



robinson ethylene glycol) group forming an amide bond with one of said amino groups; wherein R is lower alkyl; x = 2 or 4; m = 450-900; n = 1-3; and n and m are chosen so that the mol. weight of the conjugate minus the erythropoietin glycoprotein is 20-100 kDa. IT 134547-95-8P, 1-165-Erythropoietin (human clone λΗΕΡΟFL13 protein moiety reduced) RL: BPN (Biosynthetic preparation); PRP (Properties); THU (Therapeutic use); BIOL (Biological study); PREP (Preparation); USES (Uses) (amino acid sequence; erythropoietin derivs. for increasing bone marrow production of reticulocytes and erythrocytes) 134547-95-8 HCAPLUS RN CN 1-165-Erythropoietin (human clone AHEPOFL13 protein moiety reduced) (9CI) (CA INDEX NAME) \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\* 11096-26-7D, Erythropoietin, polyethylene glycol conjugates 221039-34-5, Erythropoietin (human) RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); PEP (Physical, engineering or chemical process); PRP (Properties); THU (Therapeutic use); BIOL (Biological study); PROC (Process); USES (Uses) (erythropoietin derivs. for increasing bone marrow production of reticulocytes and erythrocytes) RN 11096-26-7 HCAPLUS

Erythropoietin (9CI) CN (CA INDEX NAME)

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

221039-34-5 HCAPLUS RN

CN Erythropoietin (human) (9CI) (CA INDEX NAME)

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

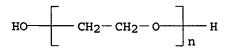
IT 25322-68-3D, Polyethylene glycol, glycoprotein

RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); PRP (Properties); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

(erythropoietin derivs. for increasing bone marrow production of reticulocytes and erythrocytes)

RN 25322-68-3 HCAPLUS

Poly(oxy-1,2-ethanediyl), α-hydro-ω-hydroxy- (9CI) (CA INDEX CN NAME)



IT 96024-34-9, Erythropoietin (human clone \( \lambda \text{HEPOFL13} \) protein moiety reduced)

RL: PRP (Properties)

(unclaimed protein sequence; erythropoietin derivs. for

increasing bone marrow production of reticulocytes and erythrocytes)

RN96024-34-9 HCAPLUS

Erythropoietin (human clone AHEPOFL13 protein moiety reduced) (9CI) CN (CA INDEX NAME)

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*



L108 ANSWER 17 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN

AN 2000:435658 HCAPLUS

DN 133:38600

- TI The use of subcutaneous **erythropoietin** and intravenous **iron** for the treatment of the anemia of severe, resistant congestive heart failure improves cardiac and renal function and functional cardiac class, and markedly reduces hospitalizations
- AU Silverberg, Donald S.; Wexler, Dov; Blum, Miriam; Keren, Gad; Sheps, David; Leibovitch, Eyal; Brosh, David; Laniado, Shlomo; Schwartz, Doron; Yachnin, Tatyana; Shapira, Itzhak; Gavish, Dov; Baruch, Ron; Koifman, Bella; Kaplan, Carl; Steinbruch, Shoshana; Iaina, Adrian
- CS Department of Nephrology and Cardiology, Tel Aviv Medical Center, Tel Aviv-Jaffa, Israel
- Journal of the American College of Cardiology (2000), 35(7),
  1737-1744
  CODEN: JACCDI; ISSN: 0735-1097
- PB Elsevier Science Inc.
- DT Journal
- LA English
- AB This study evaluated the prevalence and severity of anemia in patients with congestive heart failure (CHF) and the effect of its correction on cardiac and renal function and hospitalization. The prevalence and significance of mild anemia in patients with CHF is uncertain, and the role of erythropoietin with i.v. iron supplementation in treating this anemia is unknown. In a retrospective study, the records of the 142 patients in our CHF clinic were reviewed to find the prevalence and severity of anemia (Hb <12 g). In an intervention study, 26 of these patients, despite maximally tolerated therapy of CHF for at least six months, still had severe CHF and were also anemic. They were treated with s.c. erythropoietin and i.v. iron sufficient to increase the Hb to 12 g%. The doses of the CHF medications, except for diuretics, were not changed during the intervention period. prevalence of anemia in the 142 patients increased with the severity of CHF, reaching 79.1% in those with New York Heart Association class IV. In the intervention study, the anemia of the 26 patients was treated for a mean of 7.2±5.5 mo. The mean Hb level and mean left ventricular ejection fraction increased significantly. The mean number of hospitalizations fell by 91.9% compared with a similar period before the study. The New York Heart Association class fell significantly, as did the doses of oral and i.v. furosemide. The rate of fall of the glomerular filtration rate slowed with the treatment. Anemia is very common in CHF and its successful treatment is associated with a significant improvement in cardiac function, functional class, renal function and in a marked fall in the need for diuretics and hospitalization.
- TT 7439-89-6, Iron, biological studies
  RL: BAC (Biological activity or effector, except adverse); BPR (Biological process); BSU (Biological study, unclassified); THU (Therapeutic use);
  BIOL (Biological study); PROC (Process); USES (Uses)

(use of s.c. erythropoletin and i.v. iron for treatment of anemia of severe, resistant congestive heart failure improves cardiac and renal function and functional cardiac class, and markedly reduces hospitalizations in humans)

RN 7439-89-6 HCAPLUS

CN Iron (7CI, 8CI, 9CI) (CA INDEX NAME)

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IT 11096-26-7, Erythropoietin

RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); THU (Therapeutic use); BIOL (Biological study); USES

(use of s.c. erythropoietin and i.v. iron for treatment of anemia of severe, resistant congestive heart failure improves cardiac and renal function and functional cardiac class, and markedly reduces hospitalizations in humans) 11096-26-7 HCAPLUS

RN

CN Erythropoietin (9CI) (CA INDEX NAME)

#### \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\* RETABLE

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(RAU)	(RPY) +====-	(RVL)	(RPG)	(RWK) +====================================	File
Albitar, S	1998	13	1206	Nephrol Dial Transpl	
Anand, I	1993	70	357	Br Heart J	MEDLINE
Anand, I	1997	12	251	Curr Opin Cardiol	MEDLINE
Anon	1997	30	193	Am J Kidney Dis	
Anon	1994			Quick reference guid	
Anon	1992	20 -	32	US Renal Data System	
Besarab, A	1998	339	584	N Engl J Med	HCAPLUS
Carson, J	1995	170	32	Am J Surg	
Carson, J	1996	348	1055	Lancet	MEDLINE
Cline, C	1998	80	442	Heart	MEDLINE
Cowie, M	1997	18	208	Eur Heart J	HCAPLUS
Donnelly, S	1991	14	271	Clin Invest Med	MEDLINE
Elhalel-Dranitzki, M	1998	13	3041	Nephrol Dial Transpl	MEDLINE
Erturk, S	1999	14	1912	Nephrol Dial Transpl	
Feelders, R	1998	28	520	Eur J Clin Invest	HCAPLUS
Fishbane, S	1995	26	41	Am J Kidney Dis	MEDLINE
Foley, R	1998	9	208	J Am Soc Nephrol	
Fonarow, G	1997	30	725	J Am Coll Cardiol	MEDLINE
Ghali, J	1988	148	2013	Arch Intern Med	MEDLINE
Goch, J	1996	73	403	Nephron	MEDLINE
Goicoechea, M	1998	54	1337	Kidney Intern	HCAPLUS
Goldberg, N	1992	124	424	Am Heart J	MEDLINE
Haber, H	1991	324	353	N Engl J Med	MEDLINE
Herrera-Garza, E	1999	115	1170	Chest	MEDLINE
Hochberg, Y	1974	4	224	J Multivar Anal	
Horl, W	1999	14	50	Nephrol Dial Transpl	
King, D	1996	25	144	Age Ageing	MEDLINE
Koch, K	1995	44	201	Clin Nephrol	HCAPLUS
Kooistra, M	1998	13	828	Nephrol Dial Transpl	
Kuriyama, S	1997	77	176	Nephron	HCAPLUS
Levine, B	1990	323	236	N Engl J Med	MEDLINE
Linde, T	1996	30	115	Scand J Urol Nephrol	
Locatelli, F	1998	13	1642	Nephrol Dial Transpl	MEDLINE
Lopez-Gomez, J	1995	10	31	Nephrol Dial Transpl	
Low, I	1989	31	26	Clin Nephrol	MEDLINE
Low-Friedrich, I	1991	11	54	Am J Nephrol	MEDLINE
Ma, J	1999	10	610	J Am Soc Nephrol	MEDLINE
Macdougall, I	1996	50	1694	Kidney Int	HCAPLUS
Macdougall, I	1998	13	3030	Nephrol Dial Transpl	
Maeda, K	1982	46	137	Jpn Circ J	MEDLINE
Maschio, G	1995	10	74	Nephrol Dial Transpl	
Massie, B	1996	11	221	Curr Opin Cardiol	MEDLINE
Michaelsen, A	1998	80	437	Heart	
Opasich, C	1996	78	354	Am J Cardiol	MEDLINE
Packer, M	1999	83	1	Am J Cardiol	ĺ



Reis, S	1997	30	733	J Am Coll Cardiol	MEDLINE
Rich, M	1996	44	638	J Am Geriatr Soc	MEDLINE
Rich, M	1995	333	1190	N Engl J Med	MEDLINE
Roth, D	1994	24	777	Am J Kidney Dis	MEDLINE
Scharer, K	1993	82	953	Acta Paediatr	MEDLINE
Schwengel, R	1994	73	908	Am J Cardiol	MEDLINE
Senni, M	1997	72	453	Mayo Clin Proc	MEDLINE
Silagy, C	1993	54	84	Clin Pharmacol Ther	MEDLINE
Silverberg, D	1996	27	234	Am J Kidney Dis	HCAPLUS
Silverberg, D	1999	55	79	Kidney Int	
Silverberg, D	1996	72	413	Nephron	HCAPLUS
Silverberg, D	1998	80	1	Nephron	MEDLINE
Volpe, M	1994	74	468	Am J Cardiol	MEDLINE
Wald, M	1995	71	190	Nephron	HCAPLUS
Weil, J	1995	310	827	Br Med J	MEDLINE
Yoshida, H	1998	53	880	Kidney Int	MEDLINE

L108 ANSWER 18 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN

AN 1999:573162 HCAPLUS

DN 131:193985

TI The impact of withdrawing ACE inhibitors on **erythropoietin** responsiveness and left ventricular hypertrophy in hemodialysis patients

AU Erturk, Sehsuvar; Nergizoglu, Gokhan; Ates, Kenan; Duman, Neval; Erbay, Bulent; Karatan, Oktay; Ertug, A. Ergun

CS Department of Nephrology, Ankara University School of Medicine, Ibn-i Sina Hospital, Ankara, Turk.

SO Nephrology, Dialysis, Transplantation (1999), 14(8), 1912-1916 CODEN: NDTREA; ISSN: 0931-0509

PB Oxford University Press

DT Journal

LA English

AΒ

Background. Angiotensin-converting enzyme (ACE) inhibitors have the capability of decreasing left ventricular mass index (LVMI) in chronic hemodialysis (HD) patients. On the other hand, recent reports provide conflicting information regarding the impact of ACE inhibitors on responsiveness to recombinant human erythropoietin ( rHuEpo), and there are no data about the effect of withdrawing ACE inhibitors both on rHuEpo response and LVMI in HD patients. Methods. ACE inhibitors were switched to another antihypertensive medication in 23 out of 68 patients in our HD unit who were receiving both rHuEpo and an ACE inhibitor for more than 1 yr. Blood pressure at the pre- and post-dialysis phases, hematocrit levels and rHuEpo doses were determined at the end of the first and of the third years, and the LVMI was determined at the end of the third year. Statistical analyses were done in 15 patients in whom the study could be completed. Results. The mean (±SD) hematocrit level was increased from 26.3±6.4% to  $29.8\pm6.3$ % at the first year (P<0.05), and to  $29.4\pm6.5$ % at the third year (P<0.05 vs. before), while the mean dose of rHuEpo was decreased from 208.3±99.0 UI/kg/wk to 141.0±91.8 at the first year (P=0.01), and to 141.4 $\pm$ 81.0 at the third year (P=0.01 vs. before). Administration of rHuEpo had been stopped in two patients at the end of the first year. The mean blood pressure level and the mean LVMI were not changed (P>0.05 vs. before). There were no significant changes in dialysis parameters, iron status, plasma renin activities, and levels of aldosterone, intact parathyroid hormone, aluminum and erythropoietin. Conclusion. The findings of this small uncontrolled study indicate that withdrawal of ACE inhibitors in hypertensive chronic HD patients receiving rHuEpo may result in an increase in hematocrit level, and a decrease in dose of rHuEpo without any significant changes in the blood pressure level and LVMI.

robinson -Page 33

Controlled prospective studies are needed to clarify this issue. IT 11096-26-7, Erythropoietin

RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

(impact of withdrawing ACE inhibitors on erythropoietin responsiveness and left ventricular hypertrophy in humans on hemodialysis)

RN 11096-26-7 HCAPLUS

CN Erythropoietin (9CI) (CA INDEX NAME)

### \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\* RETABLE

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	Akpolat, T	1998	11	94	J Nephrol	MEDLINE
	Albitar, S	1998	13	1206	Nephrol Dial Transpl	HCAPLUS
	Bauer, J	1996		2331	The Kidney	
	Cannella, G	1997	30	659	Am J Kidney Dis	HCAPLUS
	Conlon, P	1994	9	1358	Nephrol Dial Transpl	MEDLINE
(	Constantinescu, C	1998	62	25	Immunol Lett	HCAPLUS
(	Cruz, D	1996	28	535	Am J Kidney Dis	HCAPLUS
I	Daugirdas, J	1993	4	1205	J Am Soc Nephrol	MEDLINE
I	Devereux, R	1986	57	450	Am J Cardiol	MEDLINE
I	Diez, J	1994	12	S31	J Hypertens	
Ι	Dyadyk, A	1997	12	945	Nephrol Dial Transpl	
ł	Erturk, S	1996	11	2050	Nephrol Dial Transpl	MEDLINE
ŀ	Erturk, S	1996	11	396	Nephrol Dial Transpl	MEDLINE
I	Escbach, J	1988	11	203	Am J Kidney Dis	
1	Escbach, J	1997	30	S192	Am J Kidney Dis	
(	Goicoechea, M	1998	54	1337	Kidney Int	HCAPLUS
(	Gossmann, J	1996	50	973	Kidney Int	HCAPLUS
(	Gould, A	1990	181	225	Eur J Pharmacol	HCAPLUS
(	Gould, A	1980	96	523	J Lab Clin Med	HCAPLUS
(	Greaves, S	1994	24	768	Am J Kidney Dis	MEDLINE
	Hirakata, H	1986	26	27	Clin Nephrol	MEDLINE
į	Julian, B	1998	9	1104	J Am Soc Nephrol	HCAPLUS
1	Kamper, A	1990	50	611	Scand J Clin Lab Inv	MEDLINE
	Macdougall, I	1994	9	1032	Nephrol Dial Transpl	
	Macdougall, I	1998	13	23	Nephrol Dial Transpl	İ
	Morrone, L	1997	64	913	Transplantation	HCAPLUS
I	Motz, W	1987	10	S148	J Cardiovasc Pharmac	
ì	Mrug, M	1997	100	2310	J Clin Invest	HCAPLUS
	Naeshiro, I	1998	354	179	Eur J Pharmacol	HCAPLUS
	Nakao, K	1967	29	754	Blood	HCAPLUS
	Navarro, J	1998	80	239	Nephron	MEDLINE
	Sahn, D	1978	58	1072	Circulation	MEDLINE
	Sasaki, M	1996	12	1403	J Hypertens	İ
	Schwenk, M	1998	18	627	Pharmacotherapy	HCAPLUS
	Sennesael, J	1985	28	A252	Kidney Int	
	Silberberg, J	1989	64	222	Am J Cardiol	MEDLINE
	Silberberg, J	1990	6	1	Can J Cardiol	MEDLINE
	Sundal, E	1991	6	955	Nephrol Dial Transpl	
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L108 ANSWER 19 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN

AΝ 1999:280686 HCAPLUS

DN 130:332154

Is there a role for adjuvant therapy in patients being treated with TIepoetin?

- AU Horl, W. H.
- CS Department of Nephrology, University of Vienna, Vienna, Austria
- SO Nephrology, Dialysis, Transplantation (1999), 14(Suppl. 2), 50-60
  CODEN: NDTREA; ISSN: 0931-0509
- PB Oxford University Press
- DT Journal; General Review
- LA English
- AB A review with 78 refs. Adjuvant therapy may allow patients being treated with epoetin to derive greater clin. benefits. Iron supplementation is currently the most widely used form of adjuvant therapy; i.v. (i.v.) iron is required by the majority of haemodialysis patients receiving epoetin. Measurement of hypochromic red blood cells is the most direct way of assessing iron supply to the bone marrow. During the correction phase, a dose of i.v. iron equivalent to 50 mg/day is recommended, with the total dose not exceeding 3 g. When subclin. vitamin C deficiency is suspected, ascorbic acid may be given orally (1-1.5 g/wk) or i.v. (300 mg three times weekly at the end of dialysis). The active vitamin D metabolites alfacalcidol and calcitriol may, under some circumstances, improve anemia and reduce epoetin dosage requirements. Vitamin B6 requirements are increased during epoetin therapy, and supplementation at a dose of 100-150 mg/wk is recommended. Supplementation of vitamin B12 is optional. Folic acid is supplemented routinely in haemodialysis patients, though evidence that it increases the efficacy of epoetin is limited. Low doses (2-3 mg/wk) should normally be sufficient to maintain optimal folic acid stores in epoetin-treated patients, although higher doses are necessary for patients with hyperhomocysteinemia. L-Carnitine supplementation may be appropriate in some patients with anemia of chronic renal failure (CRF) unresponsive to, or requiring large doses of, epoetin. Androgens potentially could reduce epoetin costs in countries with limited resources, but should only be used in men older than 50 yr with a remnant kidney. Recent animal studies indicate that the combination of epoetin and insulin-like growth factor 1 might be beneficial in CRF patients. High doses of angiotensin-converting enzyme (ACE) inhibitors should be reserved for dialysis patients who have hypertension that cannot be controlled by other agents, or who require an ACE inhibitor for treatment of heart failure.
- IT 7439-89-6, Iron, biological studies 11096-26-7
  , Epoetin

RL: THU (Therapeutic use); BIOL (Biological study); USES (Uses) (epoetin and adjuvant therapy in humans)

RN 7439-89-6 HCAPLUS

CN Iron (7CI, 8CI, 9CI) (CA INDEX NAME)

Fe

RN 11096-26-7 HCAPLUS

CN Erythropoietin (9CI) (CA INDEX NAME)

## \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

RETABLE

Referenced Author (RAU)	Year (RPY)	(RVL)	(RPG)	Referenced Work (RWK)	Referenced File
Albitar, S Albitar, S	1994 1997	9	1027	Nephrol Dial Transpl Nephrol Dial Transpl	

		LODIN	son -		
Albitar, S	1997	12	514	Nephrol Dial Transpl	HCAPLUS
Albitar, S	1998	13	1206	Nephrol Dial Transpl	
Argiles, A	1994	9	1809	Nephrol Dial Transpl	MEDLINE
Argiles, A	1994	9	1809	Nephrol Dial Transpl	MEDLINE
Azizi, M	1996	97	839	J Clin Invest	HCAPLUS
Ballal, S	1991	17	29	Am J Kidney Dis	MEDLINE
Barany, P	1997	29	565	Am J Kidney Dis	HCAPLUS
Barbour, G	1979	139	889	Arch Intern Med	MEDLINE
Berard, E	1992	62	368	Nephron	MEDLINE
Berns, J	1992	37	264	Clin Nephrol	MEDLINE
Boran, M	1996	73	314	Nephron	MEDLINE
Brox, A	1996	50	937	Kidney Int	HCAPLUS
Brox, A	1998	66	1053	Transplantation	HCAPLUS
Carozzi, S	1990	6	312	Adv Peritoneal Dial	MEDLINE
Carozzi, S	1997	43	M535	J Am Soc Artif Inter	
Christ, E	1997	82	2985	J Clin Endocrinol Me	1
Conlon, P	1994	9	1358	Nephrol Dial Transpl	MEDLINE
Consensus Group Stateme	1994	23	177	Dial Transplant	
Cruz, D	1996	28	535	Am J Kidney Dis	HCAPLUS
Descombes, E	1993	43	1319	Kidney Int	MEDLINE
Eder, M	1997	15	327	Stem Cells	HCAPLUS
Fishbane, S	1995	26	41	Am J Kidney Dis	MEDLINE
Gastaldello, K	1995	10	44	Nephrol Dial Transpl	
Gaughan, K	1997	30	495	Am J Kidney Dis	į
Gobel, V	1994	153	43	Eur J Pediatr	MEDLINE
Goicoechea, M	1998	78	23	Nephron	HCAPLUS
Gokal, R	1979	48	393	Q J Med	MEDLINE
Golper, T	1992	5	94	Semin Dial	•
Hampers, C	1967	276	551	N Engl J Med	MEDLINE
Hess, E	1996	11	749	Nephrol Dial Transpl	MEDLINE
Hunter, R	1970	1	61	Lancet	MEDLINE
Hutchison, F	1997	29	651	Am J Kidney Dis	ļ
Julian, B	1994	46	1397	Kidney Int	MEDLINE
Kamper, A	1990	50	611	Scand J Clin Lab Inv	:
Kasama, R	1996	27	680	Am J Kidney Dis	MEDLINE
Kooistra, M	1991	57	127	Nephron	MEDLINE
Kurtz, A	1990	122	323	Acta Endocrinol	HCAPLUS
Kurtz, A	1982	149	105	FEBS Lett	HCAPLUS
Kurtz, A	1988	85	7825	Proc Natl Acad Sci U	
Labonia, W	1995	26	757	Am J Kidney Dis	HCAPLUS
Labonia, W	1987	32	754	Kidney Int	HCAPLUS
Lederle, R	1990	105	1307	Dtsch Med Wochenschr	i
Macdougall, I	1989	299	157	Br Med J	MEDLINE
Macdougall, I	1992	304	225	Br Med J	MEDLINE
Macdougall, I	1996	50	1694	Kidney Int	HCAPLUS
Macdougall, I	1995	10	607	Nephrol Dial Transpl	MEDLINE
Matsumura, M	1996	72	574	Nephron	MEDLINE
Matsumura, M	1997	77	164	Nephron	HCAPLUS
Matsuzaki, Y	1996	63	33	Int J Hematol	MEDLINE
Moore, L	1992	3	105	J Renal Nutr	
Morrone, L	1997	64	913	Transplantation	HCAPLUS
Muta, K	1993	156	264	J Cell Physiol	HCAPLUS
Mydlik, M	1997	51	S56	Kidney Int	 
Ono, K	1992	38	290	Clin Nephrol	MEDLINE
Pronai, W	1995	71	395	Nephron	HCAPLUS
Rao, D	1993	328	171	Nengl J Med	MEDLINE
Rolton, H	1991	6	440	Nephrol Dial Transpl	
Sanchez, J	1995	10	1476	Nephrol Dial Transpl	1
Shimizu, T	1994	47	178	Am J Hematol	MEDLINE
Sunder-Plassmann, G	1995	10	2070	Nephrol Dial Transpl	MEDLINE

robinson -



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Taniguchi, S	1997	90	2244	Blood	HCAPLUS
Tarng, D	1998	9	227A	J Am Soc Nephrol	
Tarng, D	1998	13	2867	Nephrol Dial Transpl	HCAPLUS
Tarng, D	1997	3	S189	Nephrology	ļ
Teruel, J	1996	7	140	J Am Soc Nephrol	HCAPLUS
Teruel, J	1996	30	129	Scand J Urol Nephrol	MEDLINE
Teruel, J	1996	30	403	Scand J Urol Nephrol	HCAPLUS
Tinawi, M	1996	74	291	Nephron	MEDLINE
Urena, P	1992	7	40	Nephrol Dial Transpl	MEDLINE
Urena, P	1991	59	384	Nephron	MEDLINE
Vihervuori, E	1996	87	2075	Blood	HCAPLUS
Vlahakos, D	1991	17	199	Am J Kidney Dis	MEDLINE
Vlahakos, D	1995	43	53	Clin Nephrol	MEDLINE
Walter, J	1993	8	1428	Nephrol Dial Transpl	MEDLINE
Westwood, N	1994	86	468	Br J Haematol	HCAPLUS
Zachee, P	1992	12	188	Am J Nephrol	MEDLINE

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AN 1998:588841 HCAPLUS

DN 130:762

TI The effects of normal as compared with low hematocrit values in patients with cardiac disease who are receiving hemodialysis and **epoetin** 

AU Besarab, Anatole; Bolton, Kline; Browne, Jeffrey K.; Egrie, Joan C.; Nissenson, Allen R.; Okamoto, Douglas M.; Schwab, Steve J.; Goodkin, David

CS Division of Nephrology and Hypertension, Henry Ford Hospital, Detroit, USA

SO New England Journal of Medicine (1998), 339(9), 584-590 CODEN: NEJMAG; ISSN: 0028-4793

PB Massachusetts Medical Society

DT Journal

LA English

In patients with end-stage renal disease, anemia develops as a result of AB erythropoietin deficiency, and recombinant human erythropoietin (epoetin) is prescribed to correct the anemia partially. We examined the risks and benefits of normalizing the hematocrit in patients with cardiac disease who were undergoing hemodialysis. We studied 1233 patients with clin. evidence of congestive heart failure or ischemic heart disease who were undergoing hemodialysis: 618 patients were assigned to receive increasing doses of epoetin to achieve and maintain a hematocrit of 42 percent, and 615 were assigned to receive doses of epoetin sufficient to maintain a hematocrit of 30 percent throughout the study. The median duration of treatment was 14 mo. The primary end point was the length of time to death or a first nonfatal myocardial infarction. After 29 mo, there were 183 deaths and 19 first nonfatal myocardial infarctions among the patients in the normal-hematocrit group and 150 deaths and 14 nonfatal myocardial infarctions among those in the low-hematocrit group (risk ratio for the normal-hematocrit group as compared with the low hematocrit group, 1.3; 95 percent confidence interval, 0.9 to 1.9). Although the difference in event-free survival between the two groups did not reach the prespecified statistical stopping boundary, the study was halted. The causes of death in the two groups were similar. The mortality rates decreased with increasing hematocrit values in both groups. The patients in the normal-hematocrit group had a decline in the adequacy of dialysis and received i.v. iron dextran more often than those in the low-hematocrit group. In patients with clin. evident congestive heart failure or ischemic heart disease who are receiving hemodialysis, administration of epoetin to raise their hematocrit to 42 percent is not recommended.

IT 11096-26-7, Epoetin

RL: ADV (Adverse effect, including toxicity); BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); THU (Therapeutic use); BIOL (Biological study); USES (Uses) (effects of normal as compared with low hematocrit values in humans with cardiac disease who are receiving hemodialysis and epoetin

RN 11096-26-7 HCAPLUS

CN Erythropoietin (9CI) (CA INDEX NAME)

# \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\* RETABLE

Referenced Author (RAU)	Year (RPY)		PG (RPG)	Referenced Work (RWK)	Referenced File
=======================================	-====+	}====-		:	
Barany, P	1996	7	1472	J Am Soc Nephrol abs	
Benz, R	1996	7	1473	J Am Soc Nephrol abs	
Beusterien, K	1996	7	763	J Am Soc Nephrol	HCAPLUS
Braumann, K	1991	58	129	Nephron	MEDLINE
Canadian Erythropoietin		300	573	BMJ	
Cannella, G	1991	6	31	Nephrol Dial Transpl	MEDLINE
Collins, A	1994	5	439	J Am Soc Nephrol abs	
Collins, A	1997	8	190A	J Am Soc Nephrol abs	
Cox, D	1972	34	187	J R Stat Soc [B]	
Eschbach, J	1989	111	992	Ann Intern Med	MEDLINE
Eschbach, J	1993	4	425	J Am Soc Nephrol abs	
Eschbach, J	1993	6	180	Semin Dial	
Evans, R	1990	263	825	JAMA	MEDLINE
Goldberg, N	1992	124	424	Am Heart J	MEDLINE
Hoen, B	1995	10	377	Nephrol Dial Transpl	MEDLINE
Jennison, C	1990	5	299	Stat Sci	
Kusunoki, M	1981	1	413	J Cereb Blood Flow M	MEDLINE
Lan, K	1983	70	659	Biometrika	
Lowrie, E	1995			The anemia of ESRD a	
Lundin, A	1991	58	315	Nephron	MEDLINE
Macdougall, I	1990	335	614	Erratum Lancet	
Macdougall, I	1990	335	489	Lancet	MEDLINE
Marsh, J	1991	39	155	Kidney Int	MEDLINE
Massachusetts General H	1992	327	718	N Engl J Med	
McHorney, C	1993	31	247	Med Care	MEDLINE
Muirhead, N	1993	6	184	Semin Dial	
National Institute Of D	1996		23	Renal Data System US	
Nissenson, A	1996	7	1459	J Am Soc Nephrol abs	
Parfrey, P	1990	10	213	Am J Nephrol	MEDLINE
Parfrey, P	1991	2	2	J Am Soc Nephrol	MEDLINE
Pascual, J	1991	35	280	Clin Nephrol	MEDLINE
Rostand, S	1990		409	Clinical dialysis 2n	
Salonen, J	1992	86	803	Circulation	HCAPLUS
Sennesael, J	1991	40	121	Kidney Int	MEDLINE
Silberberg, J	1989	36	286	Kidney Int	MEDLINE
Sullivan, J	1981	1	1293	Lancet	HCAPLUS
Tielemans, C	1989	4	883	Nephrol Dial Transpl	
Veys, N	1992	19	358	Am J Kidney Dis	MEDLINE
Wizemann, V	1992	62	161	Nephron	MEDLINE
Zehnder, C	1992	61	21	Nephron	MEDLINE

L108 ANSWER 21 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN

AN 1998:329025 HCAPLUS

DN 129:67164

TI The estimation of efficacy of oral **iron** supplementation during treatment with **epoetin beta** (recombinant human



erythropoietin) in patients undergoing cardiac surgery

- ΑU Sowade, Olaf; Messinger, Diethelm; Franke, Werner; Sowade, Birgit; Scigalla, Paul; Warnke, Harry
- Department of Cardiac Surgery, Charite-Hospital, Humboldt University CS Berlin, Mannheim, Germany
- so European Journal of Haematology (1998), 60(4), 252-259 CODEN: EJHAEC; ISSN: 0902-4441
- PB Munksgaard International Publishers Ltd.
- DTJournal
- LΑ English
- We estimated the efficacy of oral iron therapy during treatment with AB rhEPO (erythropoietin) in patients undergoing cardiac surgery who were contraindicated for autologous blood donation. Seventy-six patients were enrolled in this double-blind, placebo-controlled trial and assigned to the 2 treatment groups (5+500 U/kg body weight rhEPO or placebo i.v. over 14 d before surgery). During the treatment period all patients received 300 mg Fe2+ (iron glycine sulfate) orally per day. RhEPO therapy produced significant increases in Hb concentration (Hb), reticulocyte count, hematocrit (Hct) and the hypochromic red blood cells (HRBC), and a decrease in transferrin saturation (41%) compared to the placebo group before surgery. However, the preoperative increase in HRBC was independent of the baseline ferritin and even correlated pos. with the preoperative increase in Hct (r=0.47, p<0.01). In rhEPO patients there were inverse correlations between baseline serum iron and the preoperative increases in Hb (r=-0.39, p<0.05), Hct (r=-0.50, p<0.01) and HRBC (r=-0.53, p<0.001). With this treatment regimen the HRBC appear to reflect the degree of erythropoietic stimulation rather than functional iron deficiency. The preoperative increases in reticulocytes, HRBC and Hb/Hct in patients with ferritin <100 mg/l or transferrin saturation <16% showed no significant difference compared to their complementary groups. The preoperative decrease in storage iron and the inverse correlation between the baseline ferritin and the preoperative change in ferritin (r=-0.94, p<0.0001) in the rhEPO group indicate that the iron requirement for Hb synthesis is probably covered by the breakdown of stored iron and an increase in the rate of absorption of orally administered Fe2+. I.v. rhEPO treatment with 5+500 U/kg body weight in combination with 300 mg oral Fe2+/d given over 14 d before surgery is a suitable regimen to increase Hb by about 1.61 g/dL and Hct by 0.06.
- IT 7439-89-6, Iron, biological studies
  - RL: ADV (Adverse effect, including toxicity); BIOL (Biological study) (deficiency; estimation of efficacy of oral iron supplementation during treatment with epoetin beta (recombinant
- human erythropoietin) in patients undergoing cardiac surgery)
- RN 7439-89-6 HCAPLUS
- Iron (7CI, 8CI, 9CI) CN (CA INDEX NAME)

Fe

- IT 7439-89-6D, Iron, glycine sulfate complexes, biological studies
  - RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); BIOL (Biological study)

(estimation of efficacy of oral iron supplementation during treatment with epoetin beta (recombinant human erythropoietin) in patients undergoing cardiac surgery)

- RN 7439-89-6 HCAPLUS
- CN Iron (7CI, 8CI; 9CI) (CA INDEX NAME)

Fe

### IT 11096-26-7, Erythropoietin

RL: BAC (Biological activity or effector, except adverse); BPR (Biological process); BSU (Biological study, unclassified); BIOL (Biological study); PROC (Process)

(rhEPO; estimation of efficacy of oral iron supplementation during treatment with epoetin beta (recombinant human erythropoietin) in patients undergoing cardiac surgery)

RN 11096-26-7 HCAPLUS

CN Erythropoietin (9CI) (CA INDEX NAME)

# \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\* RETABLE

Adamson, J 1993   161   Erythropoietin - mol   HCAPLUS   Beguin, Y 1997   90   A36   Blood   Blood   Biesma, D 1994   86   30   Br J Haematol   MEDLINE   Brugnara, C 1994   123   660   J Lab Clin Med   HCAPLUS   Brunner, W   1995   5   122   J Suisse Pharm   Canadian Orthopedic Per   1993   341   1227   Lancet   Cavill, I   1975   256   328   Nature   HCAPLUS   Cazzola, M   1997   89   4248   Blood   HCAPLUS   Cook, J   1990   51   301   Am J Clin Nutr   HCAPLUS   Cook, J   1990   55   68   Blood   HCAPLUS   Cook, J   1990   55   301   Am J Clin Nutr   HCAPLUS   Cook, J   1990   75   603   Br J Haematol   MEDLINE   Donohue, D   1958   37   1564   J Clin Invest   MEDLINE   Eschbach, J   1991   88   72   Contrib Nephrol   MEDLINE   Eschbach, J   1991   88   72   Contrib Nephrol   MEDLINE   Eschbach, J   1987   316   73   N Engl J Med   MEDLINE   Eschbach, J   1987   316   73   N Engl J Med   MEDLINE   Eschbach, J   1991   188   289   J Lab Clin Med   Goodnough, L   1991   188   289   J Lab Clin Med   Goodnough, L   1991   188   289   J Lab Clin Med   Horl, W   1996   11   246   Nephrol Dial Transpl   MEDLINE   Macdougall, I   1992   304   225   Br Med J   MEDLINE   Mercuriali, F   1993   33   55   Transfusion   MEDLINE   Mercuriali, F   1993   33   55   Transfusion   MEDLINE   Meyer, M   1996   129   258   J Pediatr   HCAPLUS   Nadler, S   1962   51   224   Surgery   Price, T   1996   36   29   Transfusion   MEDLINE   Silverberg, D   1996   72   413   Nephron   HCAPLUS	ced
Beguin, Y         1997         90         A36         Blood         MEDLINE           Brugnara, C         1994         86         30         Br J Haematol         MEDLINE           Brugnara, C         1994         123         660         J Lab Clin Med         HCAPLUS           Brunner, W         1995         5         122         J Suisse Pharm         HCAPLUS           Canadian Orthopedic Per 1993         341         1227         Lancet         HCAPLUS           Cavill, I         1975         256         328         Nature         HCAPLUS           Cook, J         1997         89         4248         Blood         HCAPLUS           Cook, J         1990         51         301         Am J Clin Nutr         HCAPLUS           Cook, J         1996         68         726         Blood         MEDLINE           Cook, J         1990         75         603         Br J Haematol         MEDLINE           Eschbach, J         1990         75         603         Br J Haematol         MEDLINE           Eschbach, J         1991         88         72         Contrib Nephrol         MEDLINE           Eschbach, J         1982         60         1241<	
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Brunner, W         1995         5         122         J Suisse Pharm           Canadian Orthopedic Per Cavill, I         1993         341         1227         Lancet           Cavill, I         1975         256         328         Nature         HCAPLUS           Cazzola, M         1997         89         4248         Blood         HCAPLUS           Cook, J         1990         51         301         Am J Clin Nutr         HCAPLUS           Cook, J         1986         68         726         Blood         MEDLINE           Cook, J         1990         75         603         Br J Haematol         MEDLINE           Donohue, D         1958         37         1564         J Clin Invest         MEDLINE           Eschbach, J         1991         88         72         Contrib Nephrol         MEDLINE           Eschbach, J         1987         316         73         N Engl J Med         MEDLINE           Finch, C         1982         60         1241         Blood         HCAPLUS           Goodnough, L         1991         188         289         J Lab Clin Med         MEDLINE           Horl, W         1996         11         246         Nephrol	
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Cazzola, M         1997         89         4248         Blood         HCAPLUS           Cook, J         1990         51         301         Am J Clin Nutr         HCAPLUS           Cook, J         1986         68         726         Blood         MEDLINE           Cook, J         1990         75         603         Br J Haematol         MEDLINE           Donohue, D         1958         37         1564         J Clin Invest         MEDLINE           Eschbach, J         1991         88         72         Contrib Nephrol         MEDLINE           Eschbach, J         1987         316         73         N Engl J Med         MEDLINE           Finch, C         1982         60         1241         Blood         HCAPLUS           Goodnough, L         1991         188         289         J Lab Clin Med         MEDLINE           Horl, W         1996         11         246         Nephrol Dial Transpl         MEDLINE           Kumpf, V         1990         24         162         Ann Pharmacother         MEDLINE           Mercuriali, F         1993         33         55         Transfusion         MEDLINE           Meyer, M         1996         129	
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Cook, J         1986         68         726         Blood         MEDLINE           Cook, J         1990         75         603         Br J Haematol         MEDLINE           Donohue, D         1958         37         1564         J Clin Invest         MEDLINE           Eschbach, J         1991         88         72         Contrib Nephrol         MEDLINE           Eschbach, J         1987         316         73         N Engl J Med         MEDLINE           Finch, C         1982         60         1241         Blood         HCAPLUS           Goodnough, L         1991         188         289         J Lab Clin Med         MEDLINE           Goodnough, L         1999         321         1163         N Engl J Med         MEDLINE           Horl, W         1996         11         246         Nephrol Dial Transpl         MEDLINE           Kumpf, V         1990         24         162         Ann Pharmacother         MEDLINE           Macdougall, I         1992         304         225         Br Med J         MEDLINE           Mercuriali, F         1993         33         55         Transfusion         MEDLINE           Meyer, M         1996	
Cook, J         1990         75         603         Br J Haematol         MEDLINE           Donohue, D         1958         37         1564         J Clin Invest         MEDLINE           Eschbach, J         1991         88         72         Contrib Nephrol         MEDLINE           Eschbach, J         1987         316         73         N Engl J Med         MEDLINE           Finch, C         1982         60         1241         Blood         HCAPLUS           Goodnough, L         1991         188         289         J Lab Clin Med         MEDLINE           Goodnough, L         1991         188         289         J Lab Clin Med         MEDLINE           Horl, W         1996         11         246         Nephrol Dial Transpl         MEDLINE           Kumpf, V         1990         24         162         Ann Pharmacother         MEDLINE           Macdougall, I         1992         304         225         Br Med J         MEDLINE           Mercuriali, F         1993         33         55         Transfusion         MEDLINE           Meyer, M         1996         129         258         J Pediatr         HCAPLUS           Nadler, S         1996 </td <td></td>	
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Finch, C         1982   60   1241   Blood         HCAPLUS           Goodnough, L         1991   188   289   J Lab Clin Med         MEDLINE           Goodnough, L         1989   321   1163   N Engl J Med         MEDLINE           Horl, W         1996   11   246   Nephrol Dial Transpl         MEDLINE           Kumpf, V         1990   24   162   Ann Pharmacother         MEDLINE           Macdougall, I         1992   304   225   Br Med J   MEDLINE           Mercuriali, F         1995   6   67   Erythropoiesis           Mercuriali, F         1993   33   55   Transfusion   MEDLINE           Meyer, M         1996   129   258   J Pediatr   HCAPLUS           Nadler, S         1962   51   224   Surgery           Price, T         1996   36   29   Transfusion   MEDLINE	
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Meyer, M         1996         129         258         J Pediatr         HCAPLUS           Nadler, S         1962         51         224         Surgery           Price, T         1996         36         29         Transfusion         MEDLINE	
Nadler, S 1962 51 224 Surgery Price, T 1996 36 29 Transfusion MEDLINE	
Price, T 1996 36 29 Transfusion MEDLINE	
Silverberg D 1996 72 413 Nephron LUCADING	
Skikne, B   1990   75   1870   Blood   HCAPLUS	
Skikne, B   1993     177   Erythropoietin - mol   HCAPLUS	
Skikne, B   1992   120   746   J Lab Clin Med   HCAPLUS	
Sowade, O   1997   55   89   Am J Hematol   HCAPLUS	
Sowade, O   1997   89   411   Blood   HCAPLUS	
Sowade, O   1997   129   97   J Lab Clin Med   HCAPLUS	

L108 ANSWER 22 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN

AN 1998:88081 HCAPLUS

DN 128:124107

Tİ Evaluation of erythropoietic activity on the basis of the red cell and reticulocyte distribution widths during epoetin beta therapy in patients undergoing cardiac surgery

- AU Sowade, Olaf; Sowade, Birgit; Gross, Johann; Brilla, Kay; Ziemer, Sabine; Franke, Werner; Stephan, Peter; Scigalla, Paul; Warnke, Harry
- CS Dep. Heart Surgery, Medical Fac., Humboldt Univ., Berlin, Germany
- SO Acta Haematologica (1998), 99(1), 1-7 CODEN: ACHAAH; ISSN: 0001-5792
- PB S. Karger AG
- DT Journal
- LA English
- AB The changes in the red cell and reticulocyte distribution widths during preoperative treatment with recombinant human erythropoietin (rhEPO) were evaluated in a double-blind, placebo-controlled trial in cardiac surgery patients. The increases in the reticulocyte count, in the Hb and in all distribution widths are the expression of the marked preoperative stimulation of erythropoiesis in the patients treated with Only placebo patients with a  $Hb \leq 7.5 \text{ mmol/l}$  or a transferrin > 4.0 g/l at baseline showed an increase in the red cell distribution width or in the reticulocyte Hb distribution width on oral iron therapy alone. While the reticulocyte count and the distribution widths of red cells in the rhEPO patients decreased postoperatively, only the increases in the distribution widths of reticulocytes after the second postoperative day indicate that stimulation of erythropoiesis had taken place. In patients with a low Hb or a high transferrin the rhEPO therapy should be preceded by iron therapy in order to raise the Hb level and reduce the cost of treatment.
- IT 122312-54-3, Epoetin beta

RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

(evaluation of erythropoietic activity on the basis of the red cell and reticulocyte distribution widths during **epoetin beta** therapy in patients undergoing cardiac surgery)

- RN 122312-54-3 HCAPLUS
- CN 1-165-Erythropoietin (human clone  $\lambda$ HEPOFL13 protein moiety), glycoform  $\beta$  (9CI) (CA INDEX NAME)
- \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

L108 ANSWER 23 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN

- AN 1997:466755 HCAPLUS
- DN 127:131546
- TI Kinetics of reticulocyte maturity fractions and indices and iron status during therapy with epoetin beta (recombinant human erythropoietin) in cardiac surgery patients
- AU Sowade, Olaf; Sowade, Birgit; Brilla, Kay; Franke, Werner; Stephan, Peter; Gross, Johann; Scigalla, Paul; Warnke, Harry
- CS Cardiac Surgery Clinic, Medical Faculty (Charite), Humboldt University, Berlin, Germany
- SO American Journal of Hematology (1997), 55(2), 89-96 CODEN: AJHEDD; ISSN: 0361-8609
- PB Wiley-Liss
- DT Journal
- LA English
- AB We evaluated the changes in reticulocyte maturity fractions and indexes, as measured by flow cytometry, during preoperative treatment with recombinant human erythropoietin (epoetin beta

  ) in cardiac surgery patients A total of 72 patients was enrolled in
  - ) in cardiac surgery patients. A total of 72 patients was enrolled in this double-blind, randomized, placebo-controlled clin. trial and assigned to the two treatment groups (5 + 500 U/kg bodyweight epoetin beta or placebo i.v. over 14 days preoperatively). Therapy with epoetin beta produced continuous increases in

hematocrit/Hb, in the most mature fraction of reticulocytes (LR), and in reticulocyte count. In the first treatment week there were parallel increases in the fraction of most immature reticulocytes (HR) and in the reticulocyte mean cell volume During the second week of treatment the reticulocyte mean cell Hb content (CHr) decreased, but CHr was independent of all iron parameters, affecting neither the reticulocyte fractions nor the hematocrit/Hb increase. The total preoperative rise in hematocrit correlated with the rises in LR fraction (P = 0.0270) and reticulocyte count (P = 0.0486) during the first week of treatment. Whereas in the epoetin beta patients the preoperative change in HR fraction showed neg. correlations with transferrin saturation at baseline (P = 0.0058) and with the preoperative change in iron (P = 0.0113), the preoperative change in the LR fraction correlated pos. with transferrin at baseline (P = 0.0115). Postoperatively, the reticulocyte parameters revealed that the onset of increased stimulation of erythropoiesis did not occur in the placebo patients until the second postoperative day, whereas erythropoietic activity in the epoetin beta patients was much higher during the postoperative period as well, as a result of the preoperative stimulation of erythropoiesis. reticulocyte parameters measured by flow cytometry permitted an objective anal. of erythropoietic activity during treatment with epoetin beta and in all patients post-operatively. Further studies in various types of epoetin beta therapy are needed in order to clarify the value of these reticulocyte parameters for identification of iron deficiency and optimization of epoetin beta treatment regimen.

IT 7439-89-6, Iron, biological studies

RL: BOC (Biological occurrence); BPR (Biological process); BSU (Biological study, unclassified); BIOL (Biological study); OCCU (Occurrence); PROC (Process)

(kinetics of reticulocyte maturity fractions and indexes and iron status during therapy with epoetin beta in cardiac surgery patients)

RN 7439-89-6 HCAPLUS

CN Iron (7CI, 8CI, 9CI) (CA INDEX NAME)

Fe

#### IT 122312-54-3, Epoetin beta

RL: BPR (Biological process); BSU (Biological study, unclassified); THU (Therapeutic use); BIOL (Biological study); PROC (Process); USES (Uses) (kinetics of reticulocyte maturity fractions and indexes and iron status during therapy with epoetin beta in cardiac surgery patients)

RN 122312-54-3 HCAPLUS

CN 1-165-Erythropoietin (human clone  $\lambda$ HEPOFL13 protein moiety), glycoform  $\beta$  (9CI) (CA INDEX NAME)

# \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\* RETABLE

Referenced Author (RAU)	, , ,	VOL	, , ,	Referenced Work (RWK)	Referenced File
Adamson, J Bidstrup, B Brugnara, C Brugnara, C Eschbach, J		44	725 971 623 660	Am J Med Ann Thorac Surg Am J Clin Pathol J Lab Clin Med N Engl J Med	MEDLINE MEDLINE MEDLINE HCAPLUS MEDLINE



Ganzoni, A	1969	16	1119	Br J Haematol	MEDLINE
Goodnough, L	1989	321	1163	N Engl J Med	MEDLINE
Kampf, D	1989	76	106	Contrib Nephrol	MEDLINE
Labardini, J	1973	7	301	Haematology	HCAPLUS
Lee, L	1986	7	508	Cytometry	HCAPLUS
Levine, E	1989	106	432	Surgery	MEDLINE
Lowenstein, L	1959	8	135	Int Rev Cytol	HCAPLUS
Rapoport, S	1986	1	1	The Reticulocyte	
Schmoeckel, M	1993	41	364	Thorac Cardiovasc Su	MEDLINE
Sowade, O	1995	44	257	Anaesthesist	MEDLINE
Sowade, O	1997	89	411	Blood	HCAPLUS
Sowade, O	1995	33	37a	Eur J Clin Chem Clin	
Tatsumi, N	1990	82	41	Contrib Nephrol	MEDLINE
Wells, D	1992	97	130	Am J Clin Pathol	MEDLINE
Yataganas, X	1970	62	254	Exp Cell Res	MEDLINE

L108 ANSWER 24 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN

AN 1997:51720 HCAPLUS

DN 126:127308

- TI Avoidance of allogeneic blood transfusions by treatment with epoetin beta (recombinant human erythropoietin ) in patients undergoing open-heart surgery
- AU Sowade, Olaf; Warnke, Harry; Scigalla, Paul; Sowade, Birgit; Franke, Werner; Messinger, Diethelm; Gross, Johann
- CS Med. Fac., Humboldt Univ., Berlin, Germany
- SO Blood (1997), 89(2), 411-418 CODEN: BLOOAW; ISSN: 0006-4971
- PB Saunders
- DT Journal
- LA English

AΒ

In a double-blind, randomized, placebo-controlled trial, we evaluated the ability of epoetin beta (recombinant human erythropoietin) to avoid allogeneic blood transfusions (ABT) and the associated risks in patients undergoing primary elective open-heart surgery and in whom autologous blood donation (ABD) was contraindicated. Seventy-six patients overall were enrolled onto the trial and were randomly assigned to the two treatment groups, 5+500 U/kg body weight (BW) epoetin beta or placebo i.v. over 14 days preoperatively. All patients received 300 mg Fe2+ orally per day during the treatment period. Preoperatively, the mean Hb increase was 1.50 g/dLgreater in epoetin beta patients than in placebo patients (95% confidence interval, 1.10 to 1.90 g/dL), allowing a rapid return to the baseline value by the seventh postoperative day in most epoetin beta patients. The mean volume of blood collected by intraoperative isovolemic hemodiln. was 562 mL (red blood cell mass, 274 mL) in the epoetin beta group and 218 mL (red blood cell mass, 94 mL) in the placebo group, resp. Only four patients (11%) in the epoetin beta group received an ABT, compared with 19 (53%) in the placebo group. Epoetin beta was most useful in patients with a perioperative blood loss greater than 750 mL, in those with a baseline hematocrit value less than 0.42, and in those aged ≥60 yr. The iron supplementation proved adequate despite the fact that a significant decrease in ferritin (median, 48.1%) and transferrin saturation (median, 40.5%) was observed in epoetin beta patients preoperatively. No influence of epoetin beta therapy on blood pressure, laboratory safety variables, or the frequency of specific adverse events was observed I.v. epoetin beta treatment of 5+500 U/kg BW in combination with 300 mg Fe2+ orally per day administered over 14 days preoperatively is an adequate therapy for increasing mean Hb levels by

approx. 1.50 g/dL and reducing the allogeneic blood requirement in patients undergoing elective open-heart surgery and in whom ABD is contraindicated.

IT 7439-89-6, Iron, biological studies 122312-54-3

, Epoetin beta

RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

(avoidance of allogeneic blood transfusions by treatment with epoetin beta (recombinant human erythropoietin) in combination with iron in patients

erythropoietin) in combination with iron in patients
undergoing open-heart surgery)

RN 7439-89-6 HCAPLUS

CN Iron (7CI, 8CI, 9CI) (CA INDEX NAME)

Fe

RN 122312-54-3 HCAPLUS

CN 1-165-Erythropoietin (human clone  $\lambda$ HEPOFL13 protein moiety), glycoform  $\beta$  (9CI) (CA INDEX NAME)

# \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

#### RETABLE

RETABLE					
Referenced Author	Year	VOL	PG	Referenced Work	Referenced
(RAU)	(RPY)	(RVL)	(RPG)	(RWK)	File
	h=====	-====+	<b></b>	+=====================================	+========
Bidstrup, B	1993	55	971	Ann Thorac Surg	MEDLINE
Biesma, D	1994	344	367	Lancet	MEDLINE
Bommer, J	1,988	2	406	Lancet	MEDLINE
Brugnara, C	1994	123	660	J Lab Clin Med	HCAPLUS
Canadian Orthopedic Per	1993	341	1227	Lancet	
Casagrande, J	1978	34	483	Biometrics	MEDLINE
Cosgrove, D	1985	40	380	Ann Thorac Surg	MEDLINE
Duke, M	1969	39	503	Circulation	MEDLINE
Eschbach, J	1987	316	73	N Engl J Med	MEDLINE
Geraci, J	1993	118	18	Ann Intern Med	MEDLINE
Goodnough, L	1995	60	473	Ann Thorac Surg	MEDLINE
Goodnough, L	1991.	266	86	JAMA	
Goodnough, L	1989	321	1163	N Engl J Med	MEDLINE
Hammermeister, K	1990	82	IV-380	Circulation	
Hollander, M	1973			Nonparametric Statis	
Knight, A	1988	68	681	Anesthesiology	MEDLINE
Kyo, S	1992	86	,	Circulation	
Kyo, S	1992	86	II-17	Circulation abstr 28	
Levine, E	1989	106	432	Surgery	MEDLINE
Lewis, C	1991	51	448	Ann Thorac Surg	MEDLINE
McDougall, I	1989	299	157	Br Med J	
McMahon, F	1990	76	1718	Blood	HCAPLUS
Nadler, S	1962	51	224	Surgery	
Price, T	1996	36	29	Transfusion	MEDLINE
Robertie, P	1990	28	197	Int Anaesthesiol Cli	MEDLINE
Rutherford, C	1994	96	139	Am J Med	MEDLINE
Schmoeckel, M	1993	41	364	Thorac Cardiovasc Su	MEDLINE
Schooley, J	1987	67	11	Br J Haematol	HCAPLUS
Scott, W	1992	103	1001	J Thorac Cardiovasc	MEDLINE
Skikne, B	1992	120	746	J Lab Clin Med	HCAPLUS
Sowade, O	1995	44	257	Anaesthesist	MEDLINE
Sowade, O	1995	86	352a	Blood	

Welch, H

|1992 | 116 | 393 | Ann Intern Med

MEDLINE

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L108 ANSWER 25 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN
     1996:309221 HCAPLUS
AN
DN
     125:2127
TI
     Perioperative epoetin alfa reduces transfusion
     requirements in coronary artery bypass graft surgery
ΑU
     D'Ambra, Michael
     Massachusetts General Hospital, Boston, MA, 02114, USA
CS
SO
     Seminars in Hematology (1996), 33(2, Suppl. 2), 73-74
     CODEN: SEHEA3; ISSN: 0037-1963
PB
     Saunders
DT
     Journal
LA
     English
AΒ
     Patients undergoing cardiac surgery continue to be exposed to allogeneic
            Epoetin alfa has been shown to reduce
     allogeneic blood requirements in patients scheduled for cardiac surgery,
     principally by facilitating autologous blood (AB) predonation. However,
     for some patients, there may not be sufficient time to donate AB prior to
     surgery. In this group of patients, the perisurgical use of
     epoetin alfa (recombinant human erythropoietin
     ; EPREX Janssen-Cilag, New Brunswick, NJ) warrants further
     investigation as a means of reducing allogeneic blood exposure. The
     results of an early double-blind study in 41 CABG patients suggested that
     perisurgical administration of epoetin alfa reduces
     postoperative allogeneic transfusion requirements without causing a
     significant increase in preoperative hematocrit (Hct). In a double-blind,
     placebo-controlled, parallel-group study in 182 patients scheduled for
     coronary artery bypass graft (CABG) surgery, epoetin
     alfa was administered perisurgically by s.c. (SC) injection (150
     or 300 IU/kg/d for 5 days prior to surgery, on the day of surgery, and for
     2 days postoperatively). All patients received oral iron
     supplementation for 5 days preoperatively. The intent-to-treat anal.
     showed that epoetin alfa reduced the percentage of
     patients exposed to allogeneic blood postoperatively compared with
     placebo, but not significantly. However, when the patients who developed
     surgical complications were excluded from the anal., the effect of
     epoetin alfa became significant (Fig 1).
     Reticulocytosis following surgery was significantly increased in patients
     treated with epoetin alfa 300 IU/kg. Although
     hematocrit (Hct) levels were significantly higher in epoetin
     alfa-treated patients during the first 7 days postoperatively, Hct
     levels prior to surgery were comparable among epoetin
     alfa- and placebo-treated patients. Epoetin
     alfa was well tolerated, and the overall postoperative mortality
     rate for patients treated with epoetin alfa was
     similar to that reported in other studies. Perisurgical administration of
     epoetin alfa therefore decreases exposure to allogeneic
     blood in patients undergoing CABG surgery who do not experience surgical
     complications. However, the optimum dosage regimen remains to be defined.
IT
     113427-24-0, Epoetin alfa
     RL: BAC (Biological activity or effector, except adverse); BSU (Biological
     study, unclassified); THU (Therapeutic use); BIOL (Biological study); USES
        (perioperative epoetin alfa reduces transfusion
        requirements in humans having coronary artery bypass graft surgery)
     113427-24-0 HCAPLUS
RN
CN
     1-165-Erythropoietin (human clone AHEPOFL13 protein moiety),
     glycoform \alpha (9CI) (CA INDEX NAME)
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#### \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

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L108 ANSWER 26 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN
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- AN 1996:309220 HCAPLUS
- DN 125:2126
- TI Subcutaneous **epoetin alfa** as an adjunct to autologous blood donation before elective coronary artery bypass graft surgery
- AU Gombotz, Hans
- CS Department Anesthesiology, University Graz, Graz, A-8036, Austria
- SO Seminars in Hematology (1996), 33(2, Suppl. 2), 69-72 CODEN: SEHEA3; ISSN: 0037-1963
- PB Saunders
- DT Journal
- LA English
- AΒ Autologous blood (AB) donation can minimize exposure to allogeneic blood in patients scheduled for coronary artery bypass graft (CABG) surgery. During AB donation in this group of patients, minimization of the accompanying decrease in Hb levels is important to reduce the risk of provoking silent myocardial ischemia and/or arrhythmias. Recombinant human erythropoietin (rHuEPO) has been used to facilitate AB donation and minimize the accompanying decrease in Hb levels in patients scheduled for cardiac surgery. In 24 patients scheduled for CABG surgery, once-weekly s.c. (SC) administration of rHuEPO ( epoetin alfa 400 IU/kg) plus oral iron supplementation for 4 wk prior to surgery caused marked stimulation of erythropoiesis and significantly increased collection of autologous red blood cells (RBCs) compared with oral iron alone. Furthermore, epoetin alfa minimized the decrease in Hb levels associated with AB donation and significantly attenuated allogeneic blood requirements by facilitating the collection of 4 AB units prior to surgery. During AB donation, no changes in the incidence or severity of ischemic attacks or ST-segment changes were observed using electrocardiog. monitoring. Epoetin alfa was well tolerated. Once-weekly SC administration of epoetin alfa for 4 wk
  - once-weekly SC administration of **epoetin alfa** for 4 wk therefore represents a practical means of facilitating AB donation by patients scheduled for CABG surgery.
- IT 113427-24-0, Epoetin alfa
  - RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); THU (Therapeutic use); BIOL (Biological study); USES (Uses)
    - (s.c. epoetin alfa as an adjunct to autologous blood donation before elective coronary artery bypass graft surgery in humans)
- RN 113427-24-0 HCAPLUS
- CN 1-165-Erythropoietin (human clone  $\lambda$ HEPOFL13 protein moiety), glycoform  $\alpha$  (9CI) (CA INDEX NAME)

#### \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

L108 ANSWER 27 OF 27 HCAPLUS COPYRIGHT 2005 ACS on STN

- AN 1996:309219 HCAPLUS
- DN 125:1483
- TI Autologous blood donation with recombinant human erythropoietin in cardiac surgery: The Japanese experience
- AU Baron, Jean-Francois
- CS Service d'Anesthesie Reanimation Chirurgicale, Hopital R. Broussais, Paris, 75674/14, Fr.
- SO Seminars in Hematology (1996), 33(2, Suppl. 2), 64-68 CODEN: SEHEA3; ISSN: 0037-1963
- PB Saunders



- DT Journal; General Review
- LA English
- AB A review with 9 refs. Four units of predonated autologous blood (AB) is considered sufficient to cover the blood requirements of 95% of patients undergoing elective cardiac surgery, thus avoiding the risks associated with allogeneic blood transfusion. A review of six Japanese studies was undertaken to summarize the potential for recombinant human erythropoietin (rHuEPO) to facilitate donation of AB by patients scheduled for cardiac surgery. I.v. (IV) administration of rHuEPO improved the anemia associated with AB donation, an effect that was further enhanced by IV iron supplementation. Once weekly s.c. (SC) administration of rHuEPO facilitated the donation of AB and reduced allogeneic blood requirements in patients scheduled for cardiac surgery, suggesting that rHuEPO could be administered on an outpatient basis. rhumpo was of particular benefit in anemic patients, eliminating exposure to allogeneic blood in the majority of patients. In conclusion, rHuEPO facilitates the donation of AB and reduces allogeneic blood requirements of patients scheduled for cardiac surgery.

IT 11096-26-7, Erythropoietin

RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

(autologous blood donation with recombinant human erythropoietin in Japanese cardiac surgery)

RN 11096-26-7 HCAPLUS

CN Erythropoietin (9CI) (CA INDEX NAME)

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

=> fil reg FILE 'REGISTRY' ENTERED AT 15:25:36 ON 23 AUG 2005 USE IS SUBJECT TO THE TERMS OF YOUR STN CUSTOMER AGREEMENT. PLEASE SEE "HELP USAGETERMS" FOR DETAILS. COPYRIGHT (C) 2005 American Chemical Society (ACS)

Property values tagged with IC are from the ZIC/VINITI data file provided by InfoChem.

STRUCTURE FILE UPDATES: 22 AUG 2005 HIGHEST RN 861291-85-2 DICTIONARY FILE UPDATES: 22 AUG 2005 HIGHEST RN 861291-85-2

New CAS Information Use Policies, enter HELP USAGETERMS for details.

TSCA INFORMATION NOW CURRENT THROUGH JANUARY 18, 2005

Please note that search-term pricing does apply when conducting SmartSELECT searches.

Structure search iteration limits have been increased. See HELP SLIMITS

for details.

Experimental and calculated property data are now available. For more information enter HELP PROP at an arrow prompt in the file or refer to the file summary sheet on the web at: http://www.cas.org/ONLINE/DBSS/registryss.html

=> => d ide can 1112

Carbowax 20

CN

```
L112 ANSWER 1 OF 1 REGISTRY COPYRIGHT 2005 ACS on STN
     25322-68-3 REGISTRY
RN
     Entered STN: 16 Nov 1984
ED
     Poly(oxy-1,2-ethanediyl), α-hydro-ω-hydroxy- (9CI) (CA INDEX
CN
     NAME)
OTHER NAMES:
     \alpha, \omega-Hydroxypoly(ethylene oxide)
CN
     \alpha-Hydro-\omega-hydroxypoly(oxy-1,2-ethanediyl)
CN
CN
     \alpha-Hydro-\omega-hydroxypoly (oxyethylene)
     1,2-Ethanediol, homopolymer
CN
CN
     16600
CN
     1660S
CN
     400DAB8
CN
     Alkox
     Alkox E 100
CN
CN
     Alkox E 130
CN
     Alkox E 160
CN
     Alkox E 240
CN
     Alkox E 30
     Alkox E 30G
CN
CN
     Alkox E 45
CN
     Alkox E. 60
CN
     Alkox E 75
CN
     Alkox R 100
CN
     Alkox R 1000
CN
     Alkox R 15
CN
     Alkox R 150
CN
     Alkox R 400
CN
     Alkox SR
CN
     Alkox SW
CN
     Antarox E 4000
CN
     Aquacide III
CN
     Aquaffin
CN
     Badimol
CN
     BDH 301
CN
     Bradsyn PEG
CN
     Breox 2000
CN
     Breox 20M
CN
     Breox 4000
CN
     Breox 550
CN
     Breox PEG 300
CN
     CAFO 154
CN
     Carbowax
CN
     Carbowax 100
     Carbowax 1000
CN
     Carbowax 1350
CN
     Carbowax 14000
CN
     Carbowax 1450
CN
CN
     Carbowax 1500
CN
     Carbowax 1540
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CN
     Carbowax 200
CN
     Carbowax 20000
CN
     Carbowax 25000
CN
     Carbowax 300
CN
     Carbowax 3350
ADDITIONAL NAMES NOT AVAILABLE IN THIS FORMAT - Use FCN, FIDE, or ALL for
     DISPLAY
AR
     9002-90-8
DR
     615575-04-7, 12676-74-3, 12770-93-3, 9081-95-2, 9085-02-3, 9085-03-4,
     174460-08-3, 174460-09-4, 54510-95-1, 125223-68-9, 54847-64-2, 59763-40-5,
     64441-68-5, 64640-28-4, 133573-31-6, 25104-58-9, 25609-81-8, 134919-43-0,
     101677-86-5, 99264-61-6, 106186-24-7, 112895-21-3, 114323-93-2,
     50809-04-6, 50809-59-1, 119219-06-6, 60894-12-4, 61840-14-0, 37361-15-2,
     112384-37-9, 67411-64-7, 70926-57-7, 75285-02-8, 75285-03-9, 77986-38-0,
     150872-82-5, 154394-38-4, 79964-26-4, 80341-53-3, 85399-22-0, 85945-29-5,
     90597-70-9, 88077-80-9, 88747-22-2, 34802-42-1, 107502-63-6, 107529-96-4,
     116549-90-7, 156948-19-5, 169046-53-1, 188364-77-4, 188924-03-0,
     189154-62-9, 191743-71-2, 196696-84-1, 201163-43-1, 206357-86-0,
     221638-71-7, 225502-44-3, 270910-26-4, 307928-07-0, 356055-70-4,
     391229-98-4
MF
     (C2 H4 O)n H2 O
CI
     PMS, COM
PCT Polyether
                   ADISNEWS, AGRICOLA, ANABSTR, AQUIRE, BIOBUSINESS, BIOSIS,
LC
     STN Files:
       BIOTECHNO, CA, CABA, CANCERLIT, CAPLUS, CASREACT, CBNB, CEN, CHEMCATS,
       CHEMINFORMRX, CHEMLIST, CHEMSAFE, CIN, CSCHEM, CSNB, DDFU, DETHERM*,
       DIOGENES, DRUGU, EMBASE, ENCOMPLIT, ENCOMPLIT2, ENCOMPPAT, ENCOMPPAT2, HSDB*, IFICDB, IFIPAT, IFIUDB, IPA, MEDLINE, MRCK*, MSDS-OHS, NIOSHTIC,
       PDLCOM*, PIRA, PROMT, RTECS*, SCISEARCH, SPECINFO, TOXCENTER, TULSA, ULIDAT, USAN, USPAT2, USPATFULL, VETU, VTB
          (*File contains numerically searchable property data)
     Other Sources: DSL**, TSCA**, WHO
          (**Enter CHEMLIST File for up-to-date regulatory information)
```

$$HO = \begin{bmatrix} CH_2 - CH_2 - O \end{bmatrix}_n H$$

\*\*PROPERTY DATA AVAILABLE IN THE 'PROP' FORMAT\*\*

1: 143:165633

7: 143:163291

84039 REFERENCES IN FILE CA (1907 TO DATE) 22615 REFERENCES TO NON-SPECIFIC DERIVATIVES IN FILE CA 84192 REFERENCES IN FILE CAPLUS (1907 TO DATE)

REFERENCE 2: 143:165571
REFERENCE 3: 143:165272
REFERENCE 4: 143:163889
REFERENCE 5: 143:163879
REFERENCE 6: 143:163722

REFERENCE

REFERENCE

REFERENCE 8: 143:162547

REFERENCE 9: 143:162006

REFERENCE 10: 143:161849

#### => d ide can l111 tot

L111 ANSWER 1 OF 3 REGISTRY COPYRIGHT 2005 ACS on STN

20074-52-6 REGISTRY

Entered STN: 16 Nov 1984

Iron, ion (Fe3+) (8CI, 9CI) (CA INDEX NAME) CN

OTHER NAMES:

CN Fe3+

Ferric cation CN

Ferric ion CN

Iron (Fe3+) CN

Iron ion(3+) CN

CN Iron trivalent ion

CN Iron(3+)

Iron(3+) ion CN

Iron(III) cation CN

CN Iron(III) ion

MF Fe

LC STN Files: AGRICOLA, ANABSTR, BIOBUSINESS, BIOSIS, BIOTECHNO, CA, CABA, CAPLUS, CASREACT, CHEMCATS, CHEMINFORMRX, CIN, CSNB, DDFU, DETHERM\*, DRUGU, EMBASE, IFICDB, IFIPAT, IFIUDB, NIOSHTIC, PIRA, PROMT, TOXCENTER, TULSA, ULIDAT, USPAT2, USPATFULL, VETU

(\*File contains numerically searchable property data)

Fe3+

### \*\*PROPERTY DATA AVAILABLE IN THE 'PROP' FORMAT\*\*

10606 REFERENCES IN FILE CA (1907 TO DATE) 690 REFERENCES TO NON-SPECIFIC DERIVATIVES IN FILE CA 10624 REFERENCES IN FILE CAPLUS (1907 TO DATE)

REFERENCE 1: 143:163763

REFERENCE 2: 143:160506

REFERENCE 3: 143:159768

REFERENCE 4: 143:159477

REFERENCE 5: 143:158665

REFERENCE 6: 143:158192

REFERENCE 7: 143:157906

REFERENCE 8: 143:157868

REFERENCE 9: 143:157295



#### REFERENCE 10: 143:156829

L111 ANSWER 2 OF 3 REGISTRY COPYRIGHT 2005 ACS on STN

RN 15438-31-0 REGISTRY

ED Entered STN: 16 Nov 1984

CN Iron, ion (Fe2+) (8CI, 9CI) (CA INDEX NAME)

OTHER NAMES:

CN Fe2+

CN Ferrous cation

CN Ferrous ion

CN Iron (Fe2+)

CN Iron dication

CN Iron divalent ion

CN Iron ion(2+)

CN Iron(2+)

CN Iron(II) ion

MF Fe

LC STN Files: ADISNEWS, AGRICOLA, ANABSTR, BIOBUSINESS, BIOSIS, BIOTECHNO, CA, CABA, CAPLUS, CASREACT, CEN, CHEMCATS, CHEMLIST, CIN, CSNB, DETHERM\*, EMBASE, HSDB\*, IFICDB, IFIPAT, IFIUDB, NIOSHTIC, PIRA, PROMT, RTECS\*, TOXCENTER, TULSA, ULIDAT, USPAT2, USPATFULL (\*File contains numerically searchable property data)

Fe2+

#### \*\*PROPERTY DATA AVAILABLE IN THE 'PROP' FORMAT\*\*

10171 REFERENCES IN FILE CA (1907 TO DATE)
479 REFERENCES TO NON-SPECIFIC DERIVATIVES IN FILE CA
10197 REFERENCES IN FILE CAPLUS (1907 TO DATE)

REFERENCE 1: 143:163763

REFERENCE 2: 143:162403

REFERENCE 3: 143:162235

REFERENCE 4: 143:161206

REFERENCE 5: 143:160143

REFERENCE 6: 143:158665

REFERENCE 7: 143:158598

REFERENCE 8: 143:158259

REFERENCE 9: 143:158197

REFERENCE 10: 143:157906

L111 ANSWER 3 OF 3 REGISTRY COPYRIGHT 2005 ACS on STN

RN 7439-89-6 REGISTRY

ED Entered STN: 16 Nov 1984

CN Iron (7CI, 8CI, 9CI) (CA INDEX NAME)

OTHER NAMES:

```
CN
     300A
     3ZhP
CN
     A 131
CN
CN
     A 227
     AC 325
CN
CN
     Ancor B
CN
     Ancor EN 80/150
CN
     Ancor Image 100
CN
     AQ 80
CN
     Armco 80
CN
     Armco iron
     ASC 300
CN
     ASC 300 (metal)
CN
     Atomel 300M200
CN
     Atomel 500M
CN
     Atomet 28
CN
     Atomet 95
CN
     Atomet 95G
CN
     Atomet 95SP
CN
CN
     Atomiron .44MR
CN
     Atomiron 5M
CN
     Atomiron AFP 25
CN
     Atomiron AFP 5
CN
     ATW 230
     ATW 432
CN
CN
     BASF-EW
     Carbon 0.17, iron 99.83 (atomic)
CN
     Carbonyl iron
CN
CN
     CM
     CM (iron)
CN
     Copy Powder CS 105-175
CN
CN
     DH
CN
     DKP
CN
     DKP (metal)
     DM 96
CN
CN
     DM 96 (iron)
CN
     DNK 2R
CN
     DSP 1000
     DSP 128B
CN
CN
     DSP 135
     DSP 135C
CN
     DSP 138
CN
    EF 1000
CN
     EF 250
CN
CN
     EFV
     EFV 200/300
CN
     EFV 250
CN
CN
     EFV 250/400
CN
     Electrolytic iron
CN
     EO 5A
ADDITIONAL NAMES NOT AVAILABLE IN THIS FORMAT - Use FCN, FIDE, or ALL for
     DISPLAY
     8011-79-8, 8053-60-9, 129048-51-7, 73135-38-3, 70884-35-4, 39344-71-3,
DR
     190454-13-8, 195161-83-2, 199281-22-6, 443783-52-6, 675141-17-0
MF
     Fe
CI
     COM
LC
     STN Files:
                  ADISNEWS, AGRICOLA, ANABSTR, AQUIRE, BIOBUSINESS, BIOSIS,
       BIOTECHNO, CA, CABA, CANCERLIT, CAOLD, CAPLUS, CASREACT, CBNB, CEN,
       CHEMCATS, CHEMINFORMRX, CHEMLIST, CIN, CSCHEM, CSNB, DDFU, DETHERM*,
       DIOGENES, DIPPR*, DRUGU, EMBASE, ENCOMPLIT, ENCOMPLIT2, ENCOMPPAT,
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ENCOMPPAT2, HSDB\*, IFICDB, IFIPAT, IFIUDB, IPA, MEDLINE, MRCK\*, MSDS-OHS, NIOSHTIC, PIRA, PROMT, RTECS\*, TOXCENTER, TULSA, ULIDAT, USPAT2, USPATFULL, VETU, VTB

(\*File contains numerically searchable property data)
Other Sources: DSL\*\*, EINECS\*\*, TSCA\*\*

(\*\*Enter CHEMLIST File for up-to-date regulatory information)

Fe

#### \*\*PROPERTY DATA AVAILABLE IN THE 'PROP' FORMAT\*\*

414329 REFERENCES IN FILE CA (1907 TO DATE)
21949 REFERENCES TO NON-SPECIFIC DERIVATIVES IN FILE CA
414665 REFERENCES IN FILE CAPLUS (1907 TO DATE)
1 REFERENCES IN FILE CAOLD (PRIOR TO 1967)

REFERENCE 1: 143:165644

REFERENCE 2: 143:165586

REFERENCE 3: 143:165566

REFERENCE 4: 143:165562

REFERENCE 5: 143:165540

REFERENCE 6: 143:165535

REFERENCE 7: 143:165524

REFERENCE 8: 143:165522

REFERENCE 9: 143:165278

REFERENCE 10: 143:165232

#### => d ide can l110 tot

L110 ANSWER 1 OF 15 REGISTRY COPYRIGHT 2005 ACS on STN

RN 702719-62-8 REGISTRY

ED Entered STN: 02 Jul 2004

CN Erythropoietin (human) (9CI) (CA INDEX NAME)

OTHER NAMES:

CN 2: PN: WO2004047858 SEQID: 2 claimed protein

FS PROTEIN SEQUENCE

MF Unspecified

CI MAN

SR CA

LC STN Files: CA, CAPLUS, TOXCENTER, USPATFULL

#### \*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

\*\*\* USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE \*\*\*

1 REFERENCES IN FILE CA (1907 TO DATE)

1 REFERENCES IN FILE CAPLUS (1907 TO DATE)

REFERENCE 1: 141:34188

L110 ANSWER 2 OF 15 REGISTRY COPYRIGHT 2005 ACS on STN

RN 702719-61-7 REGISTRY

ED Entered STN: 02 Jul 2004

CN Erythropoietin (human) (9CI) (CA INDEX NAME)

OTHER NAMES:

CN 1: PN: WO2004047858 SEQID: 1 claimed protein

FS PROTEIN SEQUENCE

MF Unspecified

CI MAN

SR CA

LC STN Files: CA, CAPLUS, TOXCENTER, USPATFULL

\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

\*\*\* USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE \*\*\*

1 REFERENCES IN FILE CA (1907 TO DATE)

1 REFERENCES IN FILE CAPLUS (1907 TO DATE)

REFERENCE 1: 141:34188

L110 ANSWER 3 OF 15 REGISTRY COPYRIGHT 2005 ACS on STN

RN 681860-67-3 REGISTRY

ED Entered STN: 14 May 2004

CN Erythropoietin (human 165-amino acid isoform) (9CI) (CA INDEX NAME)

OTHER NAMES:

CN 73: PN: WO2004033651 SEQID: 73 claimed protein

FS PROTEIN SEQUENCE

MF Unspecified

CI MAN

SR CA

LC STN Files: CA, CAPLUS

\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

\*\*\* USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE \*\*\*

1 REFERENCES IN FILE CA (1907 TO DATE)

1 REFERENCES TO NON-SPECIFIC DERIVATIVES IN FILE CA

1 REFERENCES IN FILE CAPLUS (1907 TO DATE)

REFERENCE 1: 140:352406

L110 ANSWER 4 OF 15 REGISTRY COPYRIGHT 2005 ACS on STN

RN 668496-69-3 REGISTRY

ED Entered STN: 29 Mar 2004

CN Erythropoietin (human 166-amino acids variant) (9CI) (CA INDEX NAME)

OTHER NAMES:

CN 2: PN: WO2004019972 SEQID: 2 claimed protein

FS PROTEIN SEQUENCE

MF Unspecified

CI MAN

SR CA

LC STN Files: CA, CAPLUS, TOXCENTER, USPATFULL



#### \*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

\*\*\* USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE \*\*\*

1 REFERENCES IN FILE CA (1907 TO DATE)

1 REFERENCES IN FILE CAPLUS (1907 TO DATE)

#### REFERENCE 1: 140:229921

L110 ANSWER 5 OF 15 REGISTRY COPYRIGHT 2005 ACS on STN

RN 668496-68-2 REGISTRY

ED Entered STN: 29 Mar 2004

CN Erythropoietin (human 165-amino acids variant) (9CI) (CA INDEX NAME)

#### OTHER NAMES:

CN 1: PN: WO2004019972 SEQID: 1 claimed protein

FS PROTEIN SEQUENCE

MF Unspecified

CI MAN

SR CA

LC STN Files: CA, CAPLUS, TOXCENTER, USPATFULL

#### \*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

\*\*\* USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE \*\*\*

1 REFERENCES IN FILE CA (1907 TO DATE)

1 REFERENCES IN FILE CAPLUS (1907 TO DATE)

#### REFERENCE 1: 140:229921

L110 ANSWER 6 OF 15 REGISTRY COPYRIGHT 2005 ACS on STN

RN **510776-48-4** REGISTRY

ED Entered STN: 06 May 2003

CN 29-165-erythropoietin (human) (9CI) (CA INDEX NAME)

OTHER NAMES:

CN 3: PN: WO03029291 SEQID: 1 claimed protein

FS PROTEIN SEQUENCE

MF Unspecified

CI MAN

SR CA

LC STN Files: CA, CAPLUS, USPAT2, USPATFULL

## \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

\*\*\* USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE \*\*\*

1 REFERENCES IN FILE CA (1907 TO DATE)

1 REFERENCES IN FILE CAPLUS (1907 TO DATE)

#### REFERENCE 1: 138:298131

L110 ANSWER 7 OF 15 REGISTRY COPYRIGHT 2005 ACS on STN

RN 510776-47-3 REGISTRY

ED Entered STN: 06 May 2003

CN Erythropoietin (human 166-amino acid isoform) (9CI) (CA INDEX NAME)

#### OTHER NAMES:

CN 2: PN: WO03029291 SEQID: 2 claimed protein

FS PROTEIN SEQUENCE

MF Unspecified

CI MAN

SR CA

STN Files: CA, CAPLUS, USPAT2, USPATFULL LC

\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

\*\*\* USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE \*\*\*

1 REFERENCES IN FILE CA (1907 TO DATE)

1 REFERENCES TO NON-SPECIFIC DERIVATIVES IN FILE CA

1 REFERENCES IN FILE CAPLUS (1907 TO DATE)

1: 138:298131 REFERENCE

L110 ANSWER 8 OF 15 REGISTRY COPYRIGHT 2005 ACS on STN

**510776-46-2** REGISTRY RN

ED Entered STN: 06 May 2003

Erythropoietin (human 165-amino acid isoform) (9CI) (CA INDEX CN NAME)

OTHER NAMES:

1: PN: WO03029291 SEQID: 1 claimed protein CN

FS PROTEIN SEQUENCE

MF Unspecified

CI MAN

SR CA

CA, CAPLUS, USPAT2, USPATFULL LC STN Files:

\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

\*\*\* USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE \*\*\*

1 REFERENCES IN FILE CA (1907 TO DATE)

1 REFERENCES TO NON-SPECIFIC DERIVATIVES IN FILE CA

1 REFERENCES IN FILE CAPLUS (1907 TO DATE)

REFERENCE 1: 138:298131

L110 ANSWER 9 OF 15 REGISTRY COPYRIGHT 2005 ACS on STN

221039-34-5 REGISTRY RN

Entered STN: 08 Apr 1999 ED

Erythropoietin (human) (9CI) (CA INDEX NAME) CN

OTHER NAMES:

1: PN: WO0027869 SEQID: 1 claimed protein CN

PROTEIN SEQUENCE FS

Unspecified MF

CI MAN

SR CA

STN Files: CA, CAPLUS, USPATFULL LC

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

\*\*\* USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE \*\*\*

5 REFERENCES IN FILE CA (1907 TO DATE)

2 REFERENCES TO NON-SPECIFIC DERIVATIVES IN FILE CA

5 REFERENCES IN FILE CAPLUS (1907 TO DATE)

REFERENCE 1: 134:91083

2: 132:352760 REFERENCE

REFERENCE 3: 132:31780

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REFERENCE
         4: 130:357233
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REFERENCE 5: 130:218746

L110 ANSWER 10 OF 15 REGISTRY COPYRIGHT 2005 ACS on STN

209810-58-2 REGISTRY

Entered STN: 12 Aug 1998 ED

Erythropoietin [30-asparagine, 32-threonine, 87-valine, 88-asparagine, 90-CN threonine] (human) (9CI) (CA INDEX NAME)

### OTHER NAMES:

- Aranesp CN
- Bone morphogenic protein 7 CN
- Darbepoetin alfa CN
- CN Darbepoetin alpha
- CN erythropoietin [30-asparagine, 32-threonine, 87-valine, 88-asparagine, 90threonine] (human)
- KRN 321 CN
- CN NESP
- CN Nespo
- CN Ostogenes protein 1
- FS PROTEIN SEQUENCE
- MF Unspecified
- CI MAN
- SR CAS Client Services
- STN Files: ADISINSIGHT, BIOSIS, CA, CANCERLIT, CAPLUS, IMSDRUGNEWS, LC IMSPATENTS, IMSRESEARCH, IPA, MEDLINE, MRCK\*, PATDPASPC, PHAR, PROUSDDR, RTECS\*, TOXCENTER, USAN, USPAT2, USPATFULL (\*File contains numerically searchable property data)

#### \*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

- \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*
- \*\*\* USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE \*\*\*

#### \*\*PROPERTY DATA AVAILABLE IN THE 'PROP' FORMAT\*\*

- 134 REFERENCES IN FILE CA (1907 TO DATE)
- 2 REFERENCES TO NON-SPECIFIC DERIVATIVES IN FILE CA
- 134 REFERENCES IN FILE CAPLUS (1907 TO DATE)
- REFERENCE 1: 143:133095
- REFERENCE 2: 143:110178
- REFERENCE 3: 143:89919
- REFERENCE 4: 143:72269
- REFERENCE 5: 143:72173
- REFERENCE 6: 143:72169
- REFERENCE 7: 143:71862
- REFERENCE 8: 143:71764
- REFERENCE 9: 143:19957
- REFERENCE 10: 143:19576



```
L110 ANSWER 11 OF 15 REGISTRY COPYRIGHT 2005 ACS on STN
     134547-95-8 REGISTRY
RN
     Entered STN: 28 Jun 1991
ED
CN
     1-165-Erythropoietin (human clone \( \text{AHEPOFL13} \) protein moiety
     reduced) (9CI) (CA INDEX NAME)
OTHER NAMES:
     1: PN: EP1064951 SEQID: 1 claimed protein
CN
     1: PN: WO0102017 SEQID: 1 claimed protein
1: PN: WO0187329 SEQID: 1 claimed protein
2: PN: WO0130320 SEQID: 1 unclaimed protein
CN
CN
CN
CN
     4: PN: WO0032772 TABLE: 1 claimed protein
     Erythropoietin (human 165-amino acid variant)
CN
     Erythropoietin (human isoform 1)
CN
     PROTEIN SEQUENCE
FS
MF
     Unspecified
CI
     MAN
SR
     CA
LC
     STN Files:
                   CA, CAPLUS, IMSPATENTS, IMSRESEARCH, TOXCENTER, USPATFULL
**RELATED SEQUENCES AVAILABLE WITH SEQLINK**
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
*** USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE ***
                7 REFERENCES IN FILE CA (1907 TO DATE)
                7 REFERENCES IN FILE CAPLUS (1907 TO DATE)
REFERENCE
            1: 136:11065
REFERENCE
            2:
                134:331598
REFERENCE
            3: 134:105827
REFERENCE
            4: 134:91083
REFERENCE
           5: 133:38711
REFERENCE
            6: 115:199743
REFERENCE 7: 115:23689
L110 ANSWER 12 OF 15 REGISTRY COPYRIGHT 2005 ACS on STN
RN
     122312-54-3 REGISTRY
ED
     Entered STN: 25 Aug 1989
     1-165-Erythropoietin (human clone AHEPOFL13 protein moiety),
     glycoform \beta (9CI) (CA INDEX NAME)
OTHER NAMES:
CN
     BM 06.019
CN
     EPOCH
CN
     Epoetin beta
CN
     Epogin
CN
     Marogen
CN
     NeoRecormon
CN
     NeoRecormon Multidose Vials
CN
     Recormon
FS
     PROTEIN SEQUENCE
MF
     Unspecified
CI
     MAN
SR
     US Adopted Names Council (USAN)
LC
                   ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, BIOBUSINESS,
     STN Files:
       BIOSIS, BIOTECHNO, CA, CANCERLIT, CAPLUS, CBNB, CIN, CSNB, DDFU,
```

DIOGENES, DRUGU, EMBASE, IMSCOSEARCH, IMSDRUGNEWS, IMSPATENTS, IMSRESEARCH, IPA, MEDLINE, MRCK\*, PIRA, PROMT, PROUSDDR, RTECS\*, SCISEARCH, TOXCENTER, USAN, USPAT2, USPATFULL

(\*File contains numerically searchable property data)
Other Sources: WHO

\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

\*\*\* USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE \*\*\*

\*\*PROPERTY DATA AVAILABLE IN THE 'PROP' FORMAT\*\*

101 REFERENCES IN FILE CA (1907 TO DATE)
101 REFERENCES IN FILE CAPLUS (1907 TO DATE)

REFERENCE 1: 143:120713

REFERENCE 2: 143:72269

REFERENCE 3: 143:1642

REFERENCE 4: 142:476541

REFERENCE 5: 142:404682

REFERENCE 6: 142:397729

REFERENCE 7: 142:367117

REFERENCE 8: 142:148256

REFERENCE 9: 142:107754

REFERENCE 10: 142:16877

L110 ANSWER 13 OF 15 REGISTRY COPYRIGHT 2005 ACS on STN

RN 113427-24-0 REGISTRY

ED Entered STN: 19 Mar 1988

CN 1-165-Erythropoietin (human clone  $\lambda$ HEPOFL13 protein moiety), glycoform  $\alpha$  (9CI) (CA INDEX NAME)

OTHER NAMES:

CN EPO

CN Epoade

CN Epoetin alfa

CN Epogen

CN Eprex

CN Erypo

CN Erypo 4000

CN Espo

CN Procrit

FS PROTEIN SEQUENCE

MF Unspecified

CI MAN

SR US Adopted Names Council (USAN)

LC STN Files: ADISINSIGHT, ADISNEWS, AGRICOLA, BIOBUSINESS, BIOSIS,
BIOTECHNO, CA, CANCERLIT, CAPLUS, CBNB, CEN, CIN, DIOGENES, EMBASE,
IMSCOSEARCH, IMSDRUGNEWS, IMSPATENTS, IMSRESEARCH, IPA, MEDLINE, MRCK\*,
PATDPASPC, PHAR, PIRA, PROMT, RTECS\*, TOXCENTER, USAN, USPAT2, USPATFULL
(\*File contains numerically searchable property data)

Other Sources: WHO

\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

\*\*\* USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE \*\*\*

\*\*PROPERTY DATA AVAILABLE IN THE 'PROP' FORMAT\*\*

306 REFERENCES IN FILE CA (1907 TO DATE)

4 REFERENCES TO NON-SPECIFIC DERIVATIVES IN FILE CA

306 REFERENCES IN FILE CAPLUS (1907 TO DATE)

REFERENCE 1: 143:133095

REFERENCE 2: 143:127065

REFERENCE 3: 143:127064

REFERENCE 4: 143:120713

REFERENCE 5: 143:72269

REFERENCE 6: 143:72188

REFERENCE 7: 143:71764

REFERENCE 8: 143:53890

REFERENCE 9: 143:53889

REFERENCE 10: 143:53882

L110 ANSWER 14 OF 15 REGISTRY COPYRIGHT 2005 ACS on STN

RN 96024-34-9 REGISTRY

ED Entered STN: 28 Apr 1985

CN Erythropoietin (human clone λHEPOFL13 protein moiety reduced)

(9CI) (CA INDEX NAME)

OTHER NAMES:

CN 2: PN: EP1064951 SEQID: 2 unclaimed protein

CN 2: PN: WO0102017 SEQID: 2 claimed protein

CN 2: PN: WO0136489 SEQID: 2 claimed protein

CN 2: PN: WO0187329 SEQID: 2 claimed protein

CN Erythropoietin (human 166-amino acid variant)

FS PROTEIN SEQUENCE

MF Unspecified

CI MAN

LC STN Files: CA, CAPLUS, IMSPATENTS, IMSRESEARCH, TOXCENTER, USPATFULL

\*\*RELATED SEQUENCES AVAILABLE WITH SEOLINK\*\*

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

\*\*\* USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE \*\*\*

24 REFERENCES IN FILE CA (1907 TO DATE)

4 REFERENCES TO NON-SPECIFIC DERIVATIVES IN FILE CA

24 REFERENCES IN FILE CAPLUS (1907 TO DATE)

REFERENCE 1: 136:11065

REFERENCE 2: 135:4473

REFERENCE 3: 134:105827 REFERENCE 4: 134:91083 REFERENCE 5: 133:38711 6: 131:139951 REFERENCE REFERENCE 7: 131:73971 REFERENCE 8: 129:27012 REFERENCE 9: 128:320571 REFERENCE 10: 125:50111 L110 ANSWER 15 OF 15 REGISTRY COPYRIGHT 2005 ACS on STN 11096-26-7 REGISTRY RN ED Entered STN: 16 Nov 1984 Erythropoietin (9CI) (CA INDEX NAME) CNOTHER NAMES: CNΕp CN EPO CN Epoetin Epogis S CN Hempoietine CN Unspecified MF CI COM, MAN ADISNEWS, AGRICOLA, ANABSTR, BIOBUSINESS, BIOSIS, BIOTECHNO, STN Files: LC CA, CABA, CANCERLIT, CAPLUS, CBNB, CEN, CHEMCATS, CHEMLIST, CIN, CSCHEM, CSNB, DDFU, DIOGENES, DRUGU, EMBASE, IFICDB, IFIPAT, IFIUDB, IMSPATENTS, IMSRESEARCH, IPA, MEDLINE, MRCK\*, NIOSHTIC, PHAR, PROMT, PROUSDDR, RTECS\*, SCISEARCH, TOXCENTER, USPAT2, USPATFULL, VETU (\*File contains numerically searchable property data) Other Sources: EINECS\*\* (\*\*Enter CHEMLIST File for up-to-date regulatory information) \*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\* \*\*PROPERTY DATA AVAILABLE IN THE 'PROP' FORMAT\*\* 9386 REFERENCES IN FILE CA (1907 TO DATE) 286 REFERENCES TO NON-SPECIFIC DERIVATIVES IN FILE CA 9410 REFERENCES IN FILE CAPLUS (1907 TO DATE) 1: 143:152011 REFERENCE REFERENCE 2: 143:151887 3: 143:151885 REFERENCE REFERENCE 4: 143:151210 REFERENCE 5: 143:150640

6: 143:150628

7: 143:147793

REFERENCE

REFERENCE

REFERENCE 8: 143:147789

REFERENCE 9: 143:147736

REFERENCE 10: 143:147566

=> => fil wpix

FILE 'WPIX' ENTERED AT 15:36:08 ON 23 AUG 2005 COPYRIGHT (C) 2005 THE THOMSON CORPORATION

FILE LAST UPDATED: 18 AUG 2005 <20050818/UP>
MOST RECENT DERWENT UPDATE: 200553 <200553/DW>
DERWENT WORLD PATENTS INDEX SUBSCRIBER FILE, COVERS 1963 TO DATE

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- >>> FOR INFORMATION ON ALL DERWENT WORLD PATENTS INDEX USER GUIDES, PLEASE VISIT: http://thomsonderwent.com/support/userguides/
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  FIRST VIEW FILE WPIFV.
  FOR FURTHER DETAILS: http://www.thomsonderwent.com/dwpifv <<<
- >>> THE CPI AND EPI MANUAL CODES HAVE BEEN REVISED FROM UPDATE 200501.
  PLEASE CHECK:
- => d all abeq tech abex tot 1131
- L131 ANSWER 1 OF 5 WPIX COPYRIGHT 2005 THE THOMSON CORP on STN

AN 2002-566640 [60] WPIX

DNC C2002-160608

- TI Novel conjugate of **erythropoietin** glycoprotein with polyethylene glycol, useful for treating diseases correlated with anemia in chronic renal failure patients and acquired immunodeficiency syndrome.
- DC A25 A96 B04 D16
- IN BURG, J; ENGEL, A; FRANZE, R; HILGER, B; SCHURIG, H E; TISCHER, W; WOZNY, M; BURGERT, J; SHOKOUFANDEH, R
- PA (HOFF) HOFFMANN LA ROCHE & CO AG F; (BURG-I) BURG J; (ENGE-I) ENGEL A; (FRAN-I) FRANZE R; (HILG-I) HILGER B; (SCHU-I) SCHURIG H E; (TISC-I) TISCHER W; (WOZN-I) WOZNY M

CYC 100

PI WO 2002049673 A2 20020627 (200260) \* EN 40 A61K047-48 <-RW: AT BE CH CY DE DK EA ES FI FR GB GH GM GR IE IT KE LS LU MC MW MZ

NL OA PT SD SE SL SZ TR TZ UG ZM ZW

W: AE AG AL AM AT AU AZ BA BB BG BR BY BZ CA CH CN CO CR CU CZ DE DK DM DZ EC EE ES FI GB GD GE GH GM HR HU ID IL IN IS JP KE KG KP KR KZ LC LK LR LS LT LU LV MA MD MG MK MN MW MX MZ NO NZ OM PH PL PT RO RU SD SE SG SI SK SL TJ TM TR TT TZ UA UG UZ VN YU ZA ZM ZW

US 2002115833 A1 20020822 (200262) A61K038-22

AU 2002033230 A 20020701 (200264) A61K047-48 <--

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EP 1345628
                    A2 20030924 (200363) EN
                                                     A61K047-48
        R: AL AT BE CH CY DE DK ES FI FR GB GR IE IT LI LT LU LV MC MK NL PT
           RO SE SI TR
    KR 2003074667 A 20030919 (200409)
                                                     C07K014-505
                                                                     <--
    BR 2001016381 A 20040225 (200416)
                                                     A61K047-48
                                                                     <--
                   W 20040819 (200455)
                                               68
    JP 2004525097
                                                     C07K014-505
                                                                     <--
    MX 2003005406
                    A1 20031101 (200468)
                                                     A61K047-48
                                                                     <--
    CN 1527726
                    A 20040908 (200478)
                                                     A61K047-48
    ZA 2003004647
                    A 20041124 (200481)
                                               55
                                                     A61K000-00
ADT WO 2002049673 A2 WO 2001-EP14434 20011208; US 2002115833 A1 US 2001-14363
    20011211; AU 2002033230 A AU 2002-33230 20011208; EP 1345628 A2 EP
    2001-984811 20011208, WO 2001-EP14434 20011208; KR 2003074667 A KR
    2003-708299 20030619; BR 2001016381 A BR 2001-16381 20011208, WO
    2001-EP14434 20011208; JP 2004525097 W WO 2001-EP14434 20011208, JP
    2002-551010 20011208; MX 2003005406 A1 WO 2001-EP14434 20011208, MX
    2003-5406 20030616; CN 1527726 A CN 2001-820609 20011208; ZA 2003004647 A
    ZA 2003-4647 20030613
FDT AU 2002033230 A Based on WO 2002049673; EP 1345628 A2 Based on WO
    2002049673; BR 2001016381 A Based on WO 2002049673; JP 2004525097 W Based
    on WO 2002049673; MX 2003005406 A1 Based on WO 2002049673
PRAI EP 2000-127891
                         20001220
    ICM A61K000-00; A61K038-22; A61K047-48; C07K014-505
         A61K038-18; A61P007-06; A61P013-12; A61P031-18; A61P035-00;
         C07K001-113; C07K014-575; C12P021-02
AB
    WO 200249673 A UPAB: 20021031
    NOVELTY - A conjugate (I) comprising an erythropoietin (
    EPO) glycoprotein having an N-terminal alpha -amino group, chosen
    from human EPO (hEPO) or its analogs having sequence of hEPO
    modified by addition of 1-6 glycosylation sites or a rearrangement of a
    glycosylation site, where the glycoprotein is covalently linked to a
    poly(ethylene glycol) group, is new.
         DETAILED DESCRIPTION - (I) comprises an EPO glycoprotein
```

having an N-terminal alpha -amino group and in vivo biological activity of causing bone marrow cells to increase production of reticulocytes and red blood cells, and chosen from hEPO or its analogs which have sequence of hEPO modified by addition of 1-6 glycosylation sites or a rearrangement of a glycosylation site, where the glycoprotein is covalently linked to a poly(ethylene glycol) group, with the -CO of the poly(ethylene glycol) group forming an amide bond with N-terminal alpha -amino group. The glycoprotein is covalently linked to a poly(ethylene glycol) group of formula (A).

-CO-(CH2)x - (OCH2CH2)m-OR (A)

R = methyl;

x = 2 or 3; and

m = 450-1350, 550-1000, preferably 650-750.

INDEPENDENT CLAIMS are also included for the following:

- (1) a pharmaceutical composition (II) comprising (I);
- (2) preparing (I);
- (3) a conjugate prepared by the above method; and
- (4) EPO glycoproteins comprising a sequence of 165 or 166 amino acids defined in the specification, having a N-terminal peptidic extension which represents a proteolytic cleavage site, optionally comprising a N-terminal purification tag.

ACTIVITY - Antianemic; anti-HIV; cytostatic.

No supporting data is given.

MECHANISM OF ACTION - None given.

USE - (I) Is useful for preparing medicaments for the treatment and prophylaxis of diseases correlated with anemia in chronic renal failure patients (CRF), acquired immunodeficiency syndrome (AIDS) and for treating cancer patients undergoing chemotherapy (claimed).

robinson -

(I) Is useful for treating patients by stimulating the division and differentiation of committed erythroid progenitors in the bone marrow. ADVANTAGE - (I) Has increased circulating half-life and plasma residence time, decreased clearance, increased clinical activity in vivo, improved potency, stability, and area under the curve when compared to unmodified EPO. Dwg.0/5 FS CPI FA AB; GI; DCN MC CPI: A05-H03; A10-E01; A12-V01; B04-C03C; B04-H07; B04-H0700E; B14-F03; B14-G01; D05-H10; TECH UPTX: 20020919 TECHNOLOGY FOCUS - BIOTECHNOLOGY - Preparation: (I) Is prepared by expressing, preferably serum free fermenting recombinant EPO protein comprising a N-terminal peptidic extension which comprises a proteolytic cleavage sequence, protecting the epsilon-amino groups by citraconylation, proteolytic cleavage of the N-terminal peptidic extension, pegylating the N-terminal alpha-amino group with a compound of formula (B), deprotecting epsilon-amino group of the EPO glycoprotein, and optionally carrying out purification after each of the above steps. The recombinant EPO has a sequence of 165, 166, 174, 169 or 174 amino acids defined in the specification. R = methyl;x = 2 or 3; andm = 450-1350, 550-1000, preferably 650-750. Preferred Conjugate: (I) has the formula (C). CH3O (CH2CH2O) m-CH2CH2CH2CO-NH-P (C) m = 650-750; and P = a residue of the glycoprotein without the N-terminal alpha-amino group which forms an amide linkage with the poly(ethylene glycol) group. The glycoprotein is a hEPO expressed by endogenous gene activation. The glycoprotein has the sequence of hEPO modified by a modification chosen from 22 modifications such as Asn30Thr32, Asn51Thr53, Asn57Thr59, Asn69 and Asn69Thr71, or preferably Gln24ser87Asn88Thr90, Gln38ser87Asn88Thr90 or Gln83ser87Asn88Thr90, or by a rearrangement comprising deletion of N-linked glycosylation sites in hEPO and addition of N-linked glycosylation site at position 88 of the sequence of hEPO. ABEX UPTX: 20020919 ADMINISTRATION - Administered weekly once, at a dose of 0.01-10 mug/kg, preferably 0.1-3 mug/kg. Administration routes not specified. EXAMPLE - The wild type erythropoietin (EPO) coding fragment was obtained. The coding fragment was amplified using primers EPO-EcoRI and EPO-SalI: 5'-GAGCCTGAATTCACCACC (EPO-EcoRI) 5'-AGGTGGGTCGACCTGGTCATCTGTCCCCTG (EPO-Sall) The Polymerase Chain Reaction (PCR) fragment was digested and cloned into the multiple cloning site of the pre-digested pCI-dhfr vector fragment. Expression of EPO gene was under control of the human cytomegalovirus (CMV) immediate-early enhancer/promoter region, an optimized chimeric intron for regulated expression and SV40 late polyadenylation signal. Cloning of APPRIEGR-EPO, APP-EPO or APPGAAHY-EPO expression constructs were also performed. The mutagenized cell line Chinese Hamster ovary (CHO)/dhfr-(ATCC CRL-9096) deficient in the dhfr enzyme gene was obtained. Cells were transfected with EPO plasmids using the FuGENE6 transfection reagent. Transfected cells were selected in alpha-MEM lacking nucleosides (alpha-MEM), supplemented with 10% dialyzed fetal calf serum (FCS), and 2

mM glutamine. Single colonies were isolated by fluorescence activated cell sorting (FACS), expanded, and the culture supernatants were assayed for

production and secretion of EPO. The cells were transferred into glass spinner flasks and cultivated in a hydrogen carbonate-buffered medium in a humidified CO2 incubator. Typical serum free media was used for the inoculum preparation. After the initial growth period, the cell culture was diluted with fresh medium. After 3-5 days, the culture in the fermenter was used as inoculum for further fermentation. A batch refeed process was used, i.e. when the desired cell density was reached, 80% of the culture was harvested.

The determined harvest was centrifuged, and supernatant was filtered and collected in a second cooled vessel, and purified as described in W09635718. The solution of the modified EPO was adjusted to pH 8.5-9 and stirred. Citraconic anhydride was added slowly to the solution in aliquots, pH of 9 was maintained and stirred. Residual citraconic anhydride was removed by adding 2 M ethanolamine solution. Cleavage of the modified protected EPO was achieved by adding cleavage protease or factor Xa. The removal of protease was achieved by size exclusion chromatography. The product was collected in fractions which were pooled according to the purity as analyzed by analytical reverse phase-high pressure liquid chromatography (rpHPLC).

The pooled fractions were concentrated to 7-8 mg/ml, and the pegylation reaction was performed at a molar ratio of 1:5 at a final protein concentration of 5 mg/ml. The pegylation reagent used was a methoxy-polyethylene glycol (PEG)-SBA. The 30 kDa PEG-SBA was dissolved in 1 mM HCl. Protected EPO was added and the reaction mixture was stirred. After 2 h, the reaction was stopped by adjusting the pH to 2.5 with acid. The separation of N-terminal pegylated EPO from excess reagents, reaction byproducts and non-pegylated EPO was achieved by chromatography. The product was collected in fractions which were pooled according to their purity as determined by high performance size exclusion chromatography. The PEG-A1 EPO was then concentrated to 4.5-7.5 mg/ml and stored frozen.

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L131 ANSWER 2 OF 5 WPIX COPYRIGHT 2005 THE THOMSON CORP on STN
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AN 2002-463307 [49] WPIX

DNC C2002-131715

TI Polyethylene glycol-modified **erythropoietin** obtained by chemical modification to lysine residue at 52-position, for use in drug compositions to treat anemia, especially renal anemia.

DC A25 B04 D16

IN KAWATA, H; MACHIDA, M; MIYAMOTO, H; NAKAMURA, T; SEKIMORI, Y

PA (CHUS) CHUGAI SEIYAKU KK; (KAWA-I) KAWATA H; (MACH-I) MACHIDA M; (MIYA-I) MIYAMOTO H; (NAKA-I) NAKAMURA T; (SEKI-I) SEKIMORI Y

CYC 98

PI WO 2002032957 A1 20020425 (200249)\* JA 46 C07K014-505 <-RW: AT BE CH CY DE DK EA ES FI FR GB GH GM GR IE IT KE LS LU MC MW MZ
NL OA PT SD SE SL SZ TR TZ UG ZW

W: AE AG AL AM AT AU AZ BA BB BG BR BY BZ CA CH CN CO CR CU CZ DE DK DM DZ EC EE ES FI GB GD GE GH GM HR HU ID IL IN IS JP KE KG KP KR KZ LC LK LR LS LT LU LV MA MD MG MK MN MW MX MZ NO NZ PH PL PT RO

RU SD SE SG SI SK SL TJ TM TR TT TZ UA UG US UZ VN YU ZA ZW AU 2001090312 A. 20020429 (200255) C07K014-505 <--

EP 1333036 A1 20030806 (200353) EN C07K014-505 <-R: AL AT BE CH CY DE DK ES FI FR GB GR IE IT LI LT LU LV MC MK NL PT

RO SE SI TR
JP 2002536338 X 20040226 (200416) C07K014-505 <--

US 2004082765 A1 20040429 (200429) A61K038-24

ADT WO 2002032957 A1 WO 2001-JP8539 20010928; AU 2001090312 A AU 2001-90312 20010928; EP 1333036 A1 EP 2001-970285 20010928, WO 2001-JP8539 20010928; JP 2002536338 X WO 2001-JP8539 20010928, JP 2002-536338 20010928; US 2004082765 A1 WO 2001-JP8539 20010928, US 2003-399254 20030416



FDT AU 2001090312 A Based on WO 2002032957; EP 1333036 A1 Based on WO 2002032957; JP 2002536338 X Based on WO 2002032957

PRAI JP 2000-315421

20001016

IC ICM A61K038-24; C07K014-505

ICS A61K038-22; A61K038-32; A61K047-34; **A61K047-48**; A61P007-06; A61P043-00; C07K014-23

AB WO 200232957 A UPAB: 20020802

NOVELTY - A mono-polyethylene glycol-modified **erythropoietin** (PEG-modified **EPO**) produced by chemically modifying natural **EPO** with PEG, is new.

DETAILED DESCRIPTION - INDEPENDENT CLAIMS are also included for the following:

- (1) mono-PEG-modified **EPO** compositions containing the mono-PEG-modified **EPO** and/or a PEG-modified **EPO** of natural **EPO** with 2 or more amino acid residues modified by PEG, 1 molecule of which as determined by gel filtration column chromatography in an aqueous solvent system has an apparent molecular weight of 100-900 kDa;
- (2) **EPO** preparations with long-lasting drug effect containing the PEG-modified **EPO** as active ingredient; and
- (3) preparing the PEG-modified **EPO** compositions by reacting natural **EPO** with the succcinimidyl ester derivative of PEG.

ACTIVITY - Antianemic.

MECHANISM OF ACTION - None given in source material.

USE - The modified **EPO** is for use in drug compositions to treat anemia especially renal anemia.

ADVANTAGE - The **EPO** has enhanced and high long-lasting drug effect but without damage to its physiological activity, which is obtainable by introducing PEG into a controlled binding site at a controlled number of binding molecules. With the formulated drug compositions agents, less nursing and treatment time is needed, less pain and cost to patients too.

DESCRIPTION OF DRAWING(S) - Mapped chromatographic pattern by liquid chromatography after digestion of mono-mPEG-EPO with endoprotease Lys-C: with axes of PEG binding site fixation of PEG(1)-EPO vs. intact EPO(x)(-0.5). (Drawing includes non-English language text). Dwg.2/14

FS CPI

LD CLI

FA AB; GI; DCN

MC CPI: A10-E01; A12-V01; B04-C03C; B04-H07;

B14-F03; B14-N10; D05-H10

TECH

UPTX: 20020802

TECHNOLOGY FOCUS - ORGANIC CHEMISTRY - Preferred Modified EPO: Chemical modification with PEG is at the lysine residue at 52-position of natural EPO. A mono-PEG-modified EPO is obtained by modifying natural EPO with a PEG having a molecular weight of 5-40 kDa, one molecule of which as determined by gel filtration column chromatography in an aqueous solvent system has an apparent molecular weight of 100-900 kDa.

ABEX

UPTX: 20020802

ADMINISTRATION - Administration is non-oral, e.g. intranasal or by injection at 5-50 microg.

EXAMPLE - A 0.5-ml solution of rh erythropoietin (EPO) (2.94 mg/ml) in 0.1 mM phosphate buffer at pH 8 was stirred with methoxy polyethylene glycol (PEG)-SPA (succinimidyl propionate; molecular weight of 20 kDa; 3.97 molar ratio), at room temperature for 30 minutes. Then, 10% 0.1 M glycine solution was added to deactivate the ester. After work-up and purification by chromatography on Superdex 200 HR10/30 (RTM),



3.8 mg mono-mPEG-**EPO** and 1.6 mg di-mPEG-**EPO** were obtained.

```
L131 ANSWER 3 OF 5 WPIX COPYRIGHT 2005 THE THOMSON CORP on STN
     2002-097323 [13]
AN
                        WPIX
DNC C2002-030219
TI
     Novel erythropoietin stimulating protein modified by conjugation
     to a polyethylene glycol moiety has a longer half life than the unmodified
     form and is useful to treat hematopoietic disorders.
DC
     A25 A96 B04
     BOONE, T C; FREEMAN, A; GEGG, C V; KINSTLER, O B; BOONE, T; GEGG, C;
IN
     KINSTLER, O
PA
     (AMGE-N) AMGEN INC
CYC
PΙ
     WO 2001076640
                     A2 20011018 (200213)* EN
                                                27
                                                      A61K047-48
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            DM DZ EE ES FI GB GD GE GH GM HR HU ID IL IN IS JP KE KG KP KR KZ
            LC LK LR LS LT LU LV MA MD MG MK MN MW MX MZ NO NZ PL PT RO RU SD
            SE SG SI SK SL TJ TM TR TT TZ UA UG UZ VN YU ZA ZW
     AU 2001055256
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     EP 1267942
                     A2 20030102 (200310)
                                           EN
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            RO SE SI TR
     US 6586398
                     B1 20030701 (200345)
                                                      A61K038-00
     US 2003166566
                    A1 20030904 (200359)
                                                      A61K038-22
     JP 2003530361
                     W 20031014 (200368)
                                                58
                                                      A61K047-48
     MX 2002009896
                    A1 20030301 (200413)
                                                      A61K047-48
ADT WO 2001076640 A2 WO 2001-US11346 20010406; AU 2001055256 A AU 2001-55256
     20010406; EP 1267942 A2 EP 2001-928395 20010406, WO 2001-US11346 20010406;
     US 6586398 B1 US 2000-545335 20000407; US 2003166566 A1 Cont of US
     2000-545335 20000407, US 2003-409807 20030407; JP 2003530361 W JP
     2001-574155 20010406, WO 2001-US11346 20010406; MX 2002009896 A1 WO
     2001-US11346 20010406, MX 2002-9896 20021007
FDT AU 2001055256 A Based on WO 2001076640; EP 1267942 A2 Based on WO
     2001076640; US 2003166566 A1 Cont of US 6586398; JP 2003530361 W Based on
     WO 2001076640; MX 2002009896 Al Based on WO 2001076640
                          20000407; US 2003-409807
PRAI US 2000-545335
     ICM A61K038-00; A61K038-22; A61K047-48
     ICS A61K038-18; A61P007-06; C07K017-00; C08G063-48; C08G063-91
AB
     WO 200176640 A UPAB: 20020226
     NOVELTY - A substantially homogenous preparation of chemically modified
     novel erythropoietin stimulating protein (NESP) is new.
          ACTIVITY - anti-anemia
          MECHANISM OF ACTION - increases erythropoiesis
          USE - The chemically modified NESP is used to treat a hematopoietic
     disorder (claimed).
          ADVANTAGE - The PEGylated NESP has a longer half life and so needs to
     be administered less frequently than prior art treatment with NESP or
     rHuEPO.
          DESCRIPTION OF DRAWING(S) - Hemoglobin response of normal mice after
     single bolus injections of 30 mu g/kg 30kD mono-PEG:NESP conjugate (closed
     triangle), 20kD mono-PEG:NESP conjugate (closed square), 5-kD
     poly-PEG: NESP conjugate (closed circle) or unmodified NESP (open circle)
     5-17 days post treatment.
     Dwg.21/22
```

FS

FΑ

MC

CPI

AB; GI; DCN

CPI: A12-V01; B04-C03D; B04-N02; B14-F01

robinson -

TECH

UPTX: 20020226

TECHNOLOGY FOCUS - BIOTECHNOLOGY - Preferred preparation: The NESP is preferably chemically modified with dextran, poly(n-vinyl pyurrolidone), a polyethylene glycol (PEG), a propropylene glycol homopolymer, a polypropylene oxide/ethylene oxide co-polymer, a polyoxyethylated polyol or a polyvinyl alcohol, more preferably with PEG with a molecular weight of 2-100kD, more preferably 5-30kD. The preparation may be a mixture of mono-PEGylated and poly-PEGylated NESP and is preferably comprised of at least 95% N-terminally mono-PEGylated NESP and at most 5% unPEGylated NESP. The NESP preferably has the 165 amino acid sequence fully defined in the specification. The PEG moiety is connected to NESP through aldehydes generated in the NESP carbohydrate chains or using methoxy-PEG-NHS chemistry.

ABEX

UPTX: 20020226

ADMINISTRATION - Administration is by intraperitoneal, subcutaneous or intramuscular injection, preferably with iron to maintain increased erythropoiesis. Dosage frequency is once every 4-6 weeks. EXAMPLE - 30kD mono-PEG:NESP derived by acylation with the 30kD PEG-NHS ester, 20Kd mono-PEG:NESP or 5kD polyPEG:NESP derived by reductive alkylation with the 20kD and 5kD PEG-aldehyde, or unmodified NESP respectively were administered to normal mice as a single bolus subcutaneous dose at 30, 10 or 3 mug/kg. The erythropoietic response and duration was monitored as reticulocyte count or hemoglobin concentration over time. The data showed that all three forms induced a strong erythropoietic response with significant dose reduction, and a prolonged efficacy relative to the unmodified NESP (see figure).

L131 ANSWER 4 OF 5 WPIX COPYRIGHT 2005 THE THOMSON CORP on STN 2001-147051 [15] AN WPIX

DNC C2001-043438

Novel erythropoietin-glycoprotein conjugate useful for treating diseases correlated with anemia in chronic renal failure patients, AIDS and for treating cancer, is linked to polyethylene glycol through linker. DC

TN BURG, J; HILGER, B; JOSEL, H

(HOFF) HOFFMANN LA ROCHE & CO AG F; (HOFF) ROCHE DIAGNOSTICS GMBH PA

CYC 90

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          C07K014-505
         A61P007-06; A61P013-12; A61P031-18; A61P035-00
AB
     WO 200102017 A UPAB: 20010317
     NOVELTY - A conjugate (I) comprising, human erythropoietin
     glycoprotein (EPO) having at least one free amino group and
     having in vivo biological activity of causing bone marrow cells to
     increase the production of reticulocytes and red blood cells, or its
     analogs, covalently linked to 1-3 lower-alkoxy poly(ethylene glycol)
     groups through a linker (L), is new.
          DETAILED DESCRIPTION - (I) comprises EPO or its analog
     having primary structure of human erythropoietin modified by the
     addition of 1-6 glycosylation sites or by the rearrangement of at least
     one glycosylation site. The glycoprotein is covalently linked to 1-3
     lower-alkoxy poly(ethylene glycol) groups, through a linker of formula
     -C(0)-X-S-Y', with C(0) of the linker forming an amide bond with the free
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The average molecular weight of each (PEG) moiety is 20-40

amino groups of glycoprotein.

k = 1-10; and

(ii).

X = -(CH2)k- or -CH2(O-CH2-CH2)k-;

Y' = (CH2)2-SO2-(CH2)2, CH2C(O)NH(CH2)2 or a group of formula (i) or

kilodaltons, and molecular weight of (I) is from 51-175 kilodaltons. INDEPENDENT CLAIMS are also included for the following:

- (1) a composition (II) comprising 1-90 % of (I);
- (2) a pharmaceutical composition comprising (I) or (II);
- (3) preparing (I) or (II) by covalent linking of thiol groups to an EPO, and coupling the resulting activated EPO with PEG derivative; and
  - (4) (I) or (II) prepared by the above said method.

ACTIVITY - Antianemic; anti-HIV; cytostatic.

MECHANISM OF ACTION - Enhancer of production of reticulocytes and red blood cells.

Normal healthy mice, 7-15 weeks old, were administered subcutaneously with 0.2 ml of methoxy-PEG-maleimide coupled to EPO, unmodified EPO and buffer solution. Over a period of 4 days starting 72 hours after the administration, blood was drawn by puncture of the tail vein, diluted and stained with acridine orange staining solution for 3-10 minutes. The reticulocytes were counted. The results showed superior activity and prolonged half life of the pegylated EPO species indicated by the significantly increased amounts of reticulocytes and shift of the reticulocytes count maximum using the same dose per mouse.

USE - (I) and (II) are useful for preparation of medicaments for the treatment of prophylaxis of disease correlated with anemia in chronic renal failure patients (CRF), AIDS and for the treatment of cancer patients undergoing chemotherapy. (I) and (II) are also useful for treating the above said diseases (claimed). Dwq.0/3

FS CPI

FA AB; GI; DCN

CPI: A10-E; A10-E08A; A12-V01; A12-W11L; B04-C03C; MC

B04-H07; B04-N06; B14-A02B1; B14-F03; B14-H01B

TECH UPTX: 20010317

TECHNOLOGY FOCUS - BIOTECHNOLOGY - Preferred Compound: (I) is of formula P' - (NH-CO-X-S-Y' - (OCH2CH2) m-OR) n.

m = 450-900;

n = 1-3;

R = lower alkyl; and

P' = EPO without amino group or amino group which form an amide linkage with X.

P' is also preferably of formula F1 or F2.

X = (CH2)k (preferably CH2);

k = 1-4:

m = 550-800 (preferably 650-700);

n = 1; and

R = CH3.

The average molecular weight of PEG is 24-35, preferably 30 kilodaltons. The glycoprotein is covalently linked to 1 or 2 lower alkoxy e.g. methoxy, capped PEG moieties. EPO is expressed by endogenous gene activation. Glycoprotein of EPO is modified by N30T32, N51T53, N57T59, N69, N69T71, S68N69T71, V87N88T90, S87N88T90, S87N88G89T90, \$87N88T90T92, \$87N88T90A162, N69T71\$87N88T90, N30T32V87N88T90, N89I90T91, S87N89I90T91, N136T138, N138T140, T125 or P124T125. The glycoprotein has a sequence comprising human EPO, and a second sequence at the carboxy terminus of EPO, containing at least one glycosylation site. The second sequence comprises a sequence derived from carboxy terminal sequence of human chorionic gonadotropin. The glycoprotein has a sequence SSSSKAPPPSLPSPSRLPGPSDTPILPQ, or a sequence modified by S87N88T90 or N30T32V87N88T90. The glycoprotein has the sequence of EPO modified by rearrangement of glycosylation site, preferably deletion of any of the N-linked carbohydrate sites in EPO or addition of N-linked carbohydrate site at position 88 of EPO. The

glycoprotein has a sequence Q24S87N88T90, Q38S87N88T90, or Q83S87N88T90.

Percentage of (I) in (II) is 1-90, preferably 1-96 %.

ABEX UPTX: 20010317

SPECIFIC SEQUENCES - **EPO** comprises a sequence of 165 or 166 amino acids fully defined in the specification. The glycoprotein has a sequence SSSSKAPPPSLPSPSRLPGPSDTPILPQ (claimed).

ADMINISTRATION - 0.01-10 (preferably 0.1-1) mug/kg of (I), containing 10-1000 (preferably 50-400) mug/ml of **erythropoietin** is administered by subcutaneous or intravenous injection.

EXAMPLE - 100 mg erythropoietin glycoprotein (EPO) was activated with SATA. The resulting activated EPO carrying covalently linked blocked thiol groups was separated from by-products like N-hydroxy-succinimide or non-reacted SATA by dialysis. 380 mg methoxy-PEG-maleimide was dissolved in the solution containing 95 mg activated EPO. The resulting molar ratio between activated EPO and methoxy-PEG-maleimide in the solution was 1:4. Covalently linked blocked thiol groups of activated EPO were de-blocked by 1 M aqueous hydroxylamine solution ad 30 mM. The resulting activated EPO in the reaction mixture of the solution contained free thiol (-SH) groups. Deblocking of the thiol groups was followed immediately by coupling between the activated EPO and methoxy-PEG-maleimide for 90 minutes, and 0.2 M aqueous cysteine solution ad 2 mM was added to stop coupling. After 30 minutes excess free thiol groups of the activated EPO were blocked by addition of a 0.5 M N-methylmaleimide solution in DMSO to reach a concentration of 5 mM. After 30 minutes the resulting reaction mixture now containing pegylated EPO species was dialyzed and purified. Content and purity of tri-, di- and mono-pegylated EPO species were evaluated on Coomassie-stained SDS-PAA qels.

L131 ANSWER 5 OF 5 WPIX COPYRIGHT 2005 THE THOMSON CORP on STN 2001-135308 [14] AN WPIX DNC C2001-057629 New conjugate having modified erythropoietin glycoprotein useful for stimulating red blood cell production and for treating diseases correlated with anemia in chronic renal failure, AIDS or cancer patients. DC IN BAILON, P S; SEBASTIAN BAILON, P PA (HOFF) HOFFMANN LA ROCHE & CO AG F; (BAIL-I) BAILON P S; (HOFF) HOFFMANN LA ROCHE INC CYC 40 PΙ NO 2000003372 A 20010103 (200114)\* C07K017-10 CA 2310536 A1 20010102 (200114) C07K014-505 <---DE 10031839 A1 20010201 (200114) C07K014-505 <--GB 2353281 A 20010221 (200115) C07K017-08 JP 2001064300 A 20010313 (200118) 16 C07K014-505 <--EP 1064951 A2 20010103 (200120)B EN 16 A61K047-48 <--

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          1064951 A UPAB: 20010410 ABEQ treated as Basic
     NOVELTY - A conjugate comprising an erythropoietin (EPO
     ) glycoprotein is new. The EPO has at least one free amino group
     and has the in vivo biological activity of causing bone marrow cells to
     increase production of reticulocytes and red blood cells. The glycoprotein
     is covalently linked to polyethylene glycol groups.
```

DETAILED DESCRIPTION - A conjugate comprising an erythropoietin (EPO) glycoprotein is new. The EPO has at least one free amino group and has the in vivo biological activity of causing bone marrow cells to increase production of reticulocytes and red blood cells. The glycoprotein is covalently linked to polyethylene glycol groups.

The EPO comprises human EPO (hEPO) or its analogs, which has the sequence of hEPO modified by the addition of 1-6



glycosylation sites or a rearrangement of at least one glycosylation site. The glycoprotein is covalently linked to n polyethylene glycol groups of formula CO-(CH2)x-(OCH2CH2)m-OR (I).

R = lower alkyl;

x = 2 or 3;

m = 450-900 and

n = 1-3.

n And m are chosen so that the molecular weight of the conjugate minus the erythropoietin glycoprotein is 20-100 kilodaltons. The CO of each polyethylene glycol group forms an amide bond with one of the amino groups.

INDEPENDENT CLAIMS are also included for the following:

(1) a composition comprising conjugates, each of the conjugates comprising the erythropoietin glycoprotein described above, the percentage of conjugates (where n =1) is at least 90% and

(2) preparation of (I).

ACTIVITY - Antianemic; immunostimulant; cytostatic; nephrotropic. MECHANISM OF ACTION - Bone marrow cell stimulator; erythroid progenitor stimulator.

In separate experiments, a single dose of unmodified EPO (25 ng of EPO), PEG(SBA)-EPO mixture (10 ng of conjugate), mono- and di-pegylated EPOs (10 ng conjugate), PEG(SPA)-EPO (10 ng of conjugate) and buffer solution were administered to mice. The results showed the superior activity and the prolonged half life of the pegylated EPO species indicated by the increased amounts of reticulocytes and the shift of the reticulocytes count maximum using the same dose per mouse (10 ng), compared to a dose of 25 ng for unmodified EPO. At 96 h, the amount of reticulocytes for unmodified EPO, 30 kDa PEG(SPA)-EPO, mono-SBA-EPO, di-SBA-EPO, PEG-EPO conjugate mixture and the control buffer were 500, 1406, 1501, 926, 1338 and 697, respectively. At 144 hours, the number of reticulocytes were approx. 0, 535, 607, 665, 660 and 708, respectively.

USE - Useful for the treating or preventing diseases correlated with anemia in chronic renal failure, AIDS or cancer patients undergoing chemotherapy. The conjugate or composition is also useful for preparing medicaments for the treatment or prophylaxis of these diseases (all claimed).

ADVANTAGE - Compared to unmodified EPO and conventional PEG-EPO conjugates, the conjugates have an increased circulating half-life and plasma residence time, decreased clearance, and increased clinical activity in vivo. Dwg.0/0

AB NO 200003372 A UPAB: 20010418

NOVELTY - A conjugate comprising an erythropoietin (EPO ) glycoprotein is new. The EPO has at least one free amino group and has the in vivo biological activity of causing bone marrow cells to increase production of reticulocytes and red blood cells. The glycoprotein is covalently linked to polyethylene glycol groups.

DETAILED DESCRIPTION - A conjugate comprising an erythropoietin (EPO) glycoprotein is new. The EPO has at least one free amino group and has the in vivo biological activity of causing bone marrow cells to increase production of reticulocytes and red blood cells. The glycoprotein is covalently linked to polyethylene glycol groups.

The EPO comprises human EPO (hEPO) or its analogs, which has the sequence of hEPO modified by the addition of 1-6 glycosylation sites or a rearrangement of at least one glycosylation site.

The glycoprotein is covalently linked to n polyethylene glycol groups of formula CO-(CH2)x-(OCH2CH2)m-OR (I).

R = lower alkyl; x = 2 or 3;m = 450-900 and n = 1-3.

n And m are chosen so that the molecular weight of the conjugate minus the erythropoietin glycoprotein is 20-100 kilodaltons. The CO of each polyethylene glycol group forms an amide bond with one of the amino groups.

INDEPENDENT CLAIMS are also included for the following:

(1) a composition comprising conjugates, each of the conjugates comprising the erythropoietin glycoprotein described above, the percentage of conjugates (where n =1) is at least 90% and

(2) preparation of (I).

ACTIVITY - Antianemic; immunostimulant; cytostatic; nephrotropic. MECHANISM OF ACTION - Bone marrow cell stimulator; erythroid progenitor stimulator.

In separate experiments, a single dose of unmodified EPO (25 ng of EPO), PEG(SBA)-EPO mixture (10 ng of conjugate), mono- and di-pegylated EPOs (10 ng conjugate), PEG(SPA)-EPO (10 ng of conjugate) and buffer solution were administered to mice. The results showed the superior activity and the prolonged half life of the pegylated EPO species indicated by the increased amounts of reticulocytes and the shift of the reticulocytes count maximum using the same dose per mouse (10 ng), compared to a dose of 25 ng for unmodified EPO. At 96 h, the amount of reticulocytes for unmodified EPO, 30 kDa PEG(SPA)-EPO, mono-SBA-EPO, di-SBA-EPO, PEG-EPO conjugate mixture and the control buffer were 500, 1406, 1501, 926, 1338 and 697, respectively. At 144 hours, the number of reticulocytes were approx. 0, 535, 607, 665, 660 and 708, respectively.

USE - Useful for the treating or preventing diseases correlated with anemia in chronic renal failure, AIDS or cancer patients undergoing chemotherapy. The conjugate or composition is also useful for preparing medicaments for the treatment or prophylaxis of these diseases (all claimed).

ADVANTAGE - Compared to unmodified EPO and conventional PEG-EPO conjugates, the conjugates have an increased circulating half-life and plasma residence time, decreased clearance, and increased clinical activity in vivo.

Dwg.0/0

FS CPI

FA AB; GI; DCN

MC CPI: A10-E08B; A12-V01; B04-C03C; B04-H07;

**B14-F04**; B14-G01B

ABEQ EP 1064951 A UPAB: 20010410

> NOVELTY - A conjugate comprising an erythropoietin (EPO ) glycoprotein is new. The EPO has at least one free amino group and has the in vivo biological activity of causing bone marrow cells to increase production of reticulocytes and red blood cells. The glycoprotein is covalently linked to polyethylene glycol groups.

> DETAILED DESCRIPTION - A conjugate comprising an erythropoietin (EPO) glycoprotein is new. The EPO has at least one free amino group and has the in vivo biological activity of causing bone marrow cells to increase production of reticulocytes and red blood cells. The glycoprotein is covalently linked to polyethylene glycol groups.

> The EPO comprises human EPO (hEPO) or its analogs, which has the sequence of hEPO modified by the addition of 1-6 glycosylation sites or a rearrangement of at least one glycosylation site. The glycoprotein is covalently linked to n polyethylene glycol groups

of formula CO-(CH2)x-(OCH2CH2)m-OR (I).

R = lower alkyl;

x = 2 or 3;

m = 450-900 and

n = 1-3.

n And m are chosen so that the molecular weight of the conjugate minus the **erythropoietin** glycoprotein is 20-100 kilodaltons. The CO of each polyethylene glycol group forms an amide bond with one of the amino groups.

INDEPENDENT CLAIMS are also included for the following:

(1) a composition comprising conjugates, each of the conjugates comprising the **erythropoietin** glycoprotein described above, the percentage of conjugates (where n = 1) is at least 90% and

(2) preparation of (I).

ACTIVITY - Antianemic; immunostimulant; cytostatic; nephrotropic.
MECHANISM OF ACTION - Bone marrow cell stimulator; erythroid
progenitor stimulator.

In separate experiments, a single dose of unmodified EPO (25 ng of EPO), PEG(SBA)-EPO mixture (10 ng of conjugate), mono- and di-pegylated EPOs (10 ng conjugate), PEG(SPA)-EPO (10 ng of conjugate) and buffer solution were administered to mice. The results showed the superior activity and the prolonged half life of the pegylated EPO species indicated by the increased amounts of reticulocytes and the shift of the reticulocytes count maximum using the same dose per mouse (10 ng), compared to a dose of 25 ng for unmodified EPO. At 96 h, the amount of reticulocytes for unmodified EPO, 30 kDa PEG(SPA)-EPO, mono-SBA-EPO, di-SBA-EPO, PEG-EPO conjugate mixture and the control buffer were 500, 1406, 1501, 926, 1338 and 697, respectively. At 144 hours, the number of reticulocytes were approx. 0, 535, 607, 665, 660 and 708, respectively.

USE - Useful for the treating or preventing diseases correlated with anemia in chronic renal failure, AIDS or cancer patients undergoing chemotherapy. The conjugate or composition is also useful for preparing medicaments for the treatment or prophylaxis of these diseases (all claimed).

ADVANTAGE - Compared to unmodified **EPO** and conventional PEG-**EPO** conjugates, the conjugates have an increased circulating half-life and plasma residence time, decreased clearance, and increased clinical activity in vivo.

Dwg.0/0

TECH

UPTX: 20010410

TECHNOLOGY FOCUS - ORGANIC CHEMISTRY - Preparation: Preparation of (I) comprises condensing a polymeric compound of formula (II) with a EPO glycoprotein.

Preferred compounds: The conjugate is of formula (IA) or (IB).
P = the residue of the glycoprotein without the n amino group(s), which
form amide linkage(s) with the polyethylene glycol group(s);
R = methyl;

m = 650 - 750 and

n = 1.

The glycoprotein is preferably hEPO, where the hEPO glycoprotein is expressed by endogenous gene activation. The glycoprotein has a sequence comprising 165 amino acids defined in the specification. The glycoprotein has the hEPO sequence, which has a modification selected from the following: Asn30Thr32; Asn51Thr53; Asn57Thr59; Asn69; Asn69Thr71; Ser68Asn69Thr71; Val87Asn88Thr90; Ser87Asn88Thr90; Ser87Asn88Gly89Thr90; Ser87Asn88Thr90Thr92; Ser87Asn88Thr90Ala162; Asn69Thr71Ser87Asn88Thr90; Asn30Thr32Val87Asn88Thr90; Asn89Ile90Thr91; Ser87Asn89Ile90Thr91; Asn136Thr138; Asn138Thr140; Thr125 or Pro124Thr125.





The glycoprotein also has a sequence comprising the hEPO sequence and a second sequence at the carboxy terminus of the human erythropoietin sequence, where the second sequence contains at least one glycosylation site. The second sequence comprises a sequences derived from the carboxy terminal sequence of the human chorionic gonadotropin. The glycoprotein has a sequence selected from:

- (a) the sequence hEPO and the defined 28-amino acid sequence at the carboxy terminus of the hEPO sequence;
- (b) the sequence in (a) modified by Ser87Asn88Thr90; or
- (c) the sequence in (a) modified by Asn30Thr32Val87Asn88Thr90. The glycoprotein also has the hEPO sequence modified by a rearrangement of at least one glycosylation site, where the rearrangement comprises deletion of any of the N-linked glycosylation sites in human erythropoietin and addition of an N-linked glycosylation site at position 88 of the hEPO sequence. In particular, the hEPO has a modification selected from: Gln24Ser87Asn88Thr90; Gln38Ser87Asn88Thr90; or Gln83Ser87Asn88Thr90.

Preferred composition: The percentage of conjugates in the composition, where n = 1, is at least 92%, preferably 96%.

ABEX

UPTX: 20010410

ADMINISTRATION - The dosage is 0.01-10 (preferably 0.1-1) mug/kg administered once weekly.

EXAMPLE - Erythropoietin (EPO) - producing CHO cell line (ATCC CRL8695) was prepared. A batch re-feed process was used, i.e. when the desired cell density was reached, 80% of the culture was harvested. The remaining culture was replenished with fresh culture medium and cultivated until the next harvest. The cells were removed by centrifugation or filtration and discarded. The EPO containing supernatant was in-line filtered, collected and purified. The purification of EPO-protein was disclosed in WO96/35718. The purified EPO was subjected to pegylation with mPEG-SBA (II: R = Me; x = 0-3and m = 650-750)

To 100 mg of EPOsf (9.71 ml of a 10.3 mg/ml EPOsf stock, 5.48 micro-mol), 10 ml of 0.1 M potassium phosphate buffer (pH 7.5) containing 506 mg of 30 kDa methoxy-PEG-SBA (16.5 micro-mol) was added and mixed for 2 hours at  ${\tt room\ temperature\ (20-23 degreesC)}$  . The final protein concentration was 5 mg/ml and the protein: PEG reagent ratio was 1:3. After 2 hours, the reaction was stopped by adjusting the pH to 4.5 with glacial acetic acid and stored at -20degreesC, until ready for purification. The conjugate mixture was purified, then analyzed by SDS-PAGE, and the degree of pegylation determined. The purified conjugate mixture comprised of monoand di-PEG-EPOsf and was free of unmodified EPOsf as determined by SDS-PAGE analysis. Conjugate mixture comprised 23.4 mg or 78% of the starting material.

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(FILE 'HOME' ENTERED AT 14:19:58 ON 23 AUG 2005) SET COST OFF

FILE 'HCAPLUS' ENTERED AT 14:20:07 ON 23 AUG 2005

- 1 S US20040209802/PN OR (US2003-706701# OR EP2002-26342)/AP, PRN L1 E LEHMAN P/AU
- L2 52 S E3-E7, E9-E12
  - E LEHMANN P/AU
- 267 S E3-E6, E11-E14 L3
  - E OREDDIGER R/AU
  - E ROEDDIGER R/AU

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L4
              9 S E3, E4
                E ROEDIGER R/AU
              2 S E4
L5
                E RODIGER R/AU
              1 S E4
L6
                E RODDIGER R/AU
              2 S E4
L7
                E WALTER MATSUI/AU
              4 S E4,E5
L8
                E MATSUI R/AU
             15 S E3
L9
                E MATSUI W/AU
                SEL RN L1
     FILE 'REGISTRY' ENTERED AT 14:22:38 ON 23 AUG 2005
L10
              7 S E1-E7
              6 S L10 AND ERYTHROPOIETIN
L11
L12
              1 S L10 NOT L11
                E ERYTHROPOIETIN
L13
           1792 S E3
           1792 S L11,L13
L14
                E IRON/CN
              1 S E3
L15
                E FE/MF
L16
             30 S E3 NOT MASS
             30 S L15, L16
L17
     FILE 'HCAPLUS' ENTERED AT 14:24:56 ON 23 AUG 2005
           9810 S L14
L18
          11804 S ?ERYTHROPOIETIN?
L19
L20
            129 S DARBEPOETIN? (S) (ALPHA OR ALFA)
            135 S ?DARBEPOETIN?
L21
           6067 S EPO OR EPREX
L22
            298 S EPOETIN? (S) (ALFA OR ALPHA)
L23
            100 S EPOETIN? (S) BETA
L24
L25
             458 S EPOETIN
             42 S ARANESP
L26
L27
          14463 S L18-L26
             655 S L27 AND L17
L28
L29
            1236 S L27 AND (FE OR IRON)
L30
            1243 S L28, L29
                 E HEART DISEASE/CT
                 E E4+ALL
                 E E2+ALL
L31
          86736 S E7+OLD, NT
              29 S L30 AND L31
L32
L33
               0 S E90+OLD, NT AND L30
              47 S E92+OLD, NT AND L30
L34
              36 S L32, L34 AND (PY<=2002 OR PRY<=2002 OR AY<=2002)
L35
               1 S L32, L34 AND L1-L9
L36
               3 S L35 AND ?CONJUGAT?
L37
L38
               2 S L37 NOT 3/SC
L39
               2 S L36, L38
              33 S L35 NOT L36-L39
L40
                 SEL DN AN 6-9 13-15 19-27
              16 S L40 AND E1-E48
L41
              18 S L39, L41
L42
L43
             597 S ?RHUEPO?
             155 S L43 AND (L17 OR FE OR IRON)
L44
               3 S L44 AND L31
L45
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E HEART, DISEASE/CT
               E E3+ALL
             5 S L44 AND E92+OLD, NT
L46
L47
             5 S L45, L46
L48
             4 S L47 NOT 2005/PY
L49
            19 S L42, L48 AND L1-L9, L18-L48
L50
         14474 S L27, L43
L51
           360 S L50 AND ?CONJUGAT?
L52
           330 S L50 AND ?GLYCOSYLAT?
L53
           194 S L50 AND (PEG OR PEGYLAT?)
L54
            55 S L50 AND (POLYOXYETHYLENE OR POLYETHYLENEGLYCOL OR POLYETHYLEN
L55
             4 S L50 AND POLY() (OXYETHYLENE OR ETHYLENEGLYCOL OR ETHYLENEOXIDE
           24 S L50 AND POLY() (OXY ETHYLENE OR ETHYLENE GLYCOL OR ETHYLENE OX
L56
           237 S L50 AND (POLYOXY ETHYLENE OR POLYETHYLENE GLYCOL OR POLYETHYL
L57
           316 S L50 AND POLYOXYALKYLENE
L58
    FILE 'REGISTRY' ENTERED AT 14:42:28 ON 23 AUG 2005
L59
           1 S 25322-68-3
L60
             0 S L14 AND C2H4O
    FILE 'HCAPLUS' ENTERED AT 14:42:46 ON 23 AUG 2005
           266 S L50 AND L59
L61
L62
            986 S L51-L58,L61
L63
            804 S L62 AND (PY<=2002 OR PRY<=2002 OR AY<=2002)
L64
            32 S L63 AND (L17 OR FE OR IRON)
            30 S L64 AND L18
L65
    FILE 'REGISTRY' ENTERED AT 14:44:41 ON 23 AUG 2005
L66
            1 S L14 AND NC4/ES
             11 S L14 AND S/ELS
L67
     FILE 'HCAPLUS' ENTERED AT 14:45:59 ON 23 AUG 2005
L68
         12350 S L27 AND (PY<=2002 OR PRY<=2002 OR AY<=2002)
L69
          1036 S L68 AND (L17 OR FE OR IRON)
          1808 S L63-L65,L69
L70
          1808 S L70 OR L70
L71
L72
           500 S L71 RAN=(2001:686932,)
L73
           500 S L71 RAN=(1997:740129,2001:679500)
L74
           808 S L71 RAN=(,1997:730870)
    FILE 'REGISTRY' ENTERED AT 14:48:11 ON 23 AUG 2005
     FILE 'HCAPLUS' ENTERED AT 14:48:17 ON 23 AUG 2005
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L75
            SEL L74 1- RN : 3039 TERMS
               SET SMARTSELECT OFF
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L76
          3035 S L75
     FILE 'HCAPLUS' ENTERED AT 14:48:58 ON 23 AUG 2005
               SET SMARTSELECT ON
L77
            SEL L73 1- RN : 4980 TERMS
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4980 S L77 L78

SET SMARTSELECT OFF

FILE 'HCAPLUS' ENTERED AT 14:49:40 ON 23 AUG 2005 SET SMARTSELECT ON

FILE 'REGISTRY' ENTERED AT 14:49:13 ON 23 AUG 2005

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robinson -
L79
          SEL L72 1- RN :
                         43982 TERMS
              SET SMARTSELECT OFF
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FILE 'REGISTRY' ENTERED AT 14:50:18 ON 23 AUG 2005 L80 43982 S L79 L81 49174 S L76, L78, L80 455 S'L81 AND C2H4O L82 55 S L82 AND NC4/ES L83 1 S L83 AND S/ELS L84 54 S L83 NOT L84 L85 STR L86 50 S L86 L87 L88 STR L86 50 S L88 L89 15844 S L86 FUL L90 86 S L90 AND C2H4O L91 28 S L91 AND 1/NR NOT P/ELS SEL RN 4 6-9 22 28 L93 7 S L92 AND E1-E7 L94 58 S L91 NOT L92

35 S L94 NOT P/ELS L95

### FILE 'HCAPLUS' ENTERED AT 15:13:34 ON 23 AUG 2005

8 S L93 L97 0 S L96 AND L50 136 S L51, L52 AND L53-L58, L61 L98 110 S L98 AND (PY<=2002 OR PRY<=2002 OR AY<=2002) L100 18 S L99 AND L51 AND L52 0 S L100 AND L31 L101 6 S L98 AND L31 L102 L103 4 S L99 AND L31 6 S L102, L103 L104 18 S L100 NOT L104 L105

SEL DN AN 3 7 9 12 13 15 16 17 8 S E8-E31 AND L105 L106 SEL DN AN L48 1 4 L107 2 S L48 AND E32-E37

27 S L49, L106, L107 L108 SEL HIT RN

### FILE 'REGISTRY' ENTERED AT 15:23:50 ON 23 AUG 2005

19 S E38-E56 15 S L109 AND L14 L110 L111 3 S L109 AND L17 1 S L109 AND L59 L112

FILE 'HCAPLUS' ENTERED AT 15:25:04 ON 23 AUG 2005

FILE 'REGISTRY' ENTERED AT 15:25:36 ON 23 AUG 2005

FILE 'WPIX' ENTERED AT 15:26:21 ON 23 AUG 2005

L113 1961 S L19/BI, ABEX OR L20/BI, ABEX OR L21/BI, ABEX OR L22/BI, ABEX OR L L114 570 S (B04-H07 OR C04-H07)/MC

E ERYTHROPOIETIN/CN

7 S E3-E9

E DARBEPOETIN/CN

1 S E4, E5 L116 L117 8 S L115, L116

> SEL SDCN EDIT /SDCN /DCN

robinson		Page 79
TODITIOOII		rage / J

L118	653	S	E1-E8
L119	213	s	C07K014-505/IPC
L120	2084	S	L113, L114, L118, L119
L121			L120 AND A61K047-48/IPC
L122	29	S	L120 AND A05-H03?/MC
L123	98	S	L120 AND (B04-C03C OR C04-C03C)/MC
		E	PEG/CN
L124	2	S	E3
L125	10737	S	(RAOGM6 OR R02044)/DCN OR 2044/DRN
L126	136	S	L120 AND L125
L127	33	s	L121 AND L122,L123,L126
L128	2	S	L127 AND A61P009/IPC
L129	14	s	L127 AND (B14-F? OR C14-F? OR B12-F? OR C12-F?)/MC
L130	15	s	L128, L129
		SI	EL DN AN 6 7 9 11 12
L131	5	s	E1-E10 AND L130

FILE 'WPIX' ENTERED AT 15:36:08 ON 23 AUG 2005

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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on:

August 23, 2005, 13:52:33 ; Search time 39 Seconds (without alignments) 407.071 Million cell updates/sec

US-10-706-701-1 846 Title: Perfect score: Sequence:

1 APPRLICDSRVLERYLLEAK......SNFLRGKLKLYTGEACRTGD 165 Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

Total number of hits satisfying chosen parameters: 283416 seqs, 96216763 residues Searched:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

PIR 79:\*
1: pir1:\*
2: pir2:\*
3: pir3:\*
4: pir4:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description		erythropoietin pre	erythropoietin pre						erythropoietin - r	erythropoietin - p	1	thrombopoietin - h	thrombopoietin pre		Solute binding rec	megakaryocyte grow	probable 2-hydroxy	UDP-N-acetylpyruvo	genome polyprotein	probable sensory h	ribosomal protein	ATP-dependent heli	EGF receptor subst	ABC transporter AT	conserved hypothet	methylamine utiliz	probable copper-tr	hypothetical prote	hypothetical prote	bacterioferritin X	
SUMMARIES	Ω		ZUHU	JQ0173	184613	I46083	S28148	146401	A24902	JC7699	146578	146199	G02729	180105	AB0323	AE0959	A55530	AI0443	A83274	GNWVJ8	T35681	856639	AF0526	A54696	T35450	AG2919	H97693	S36741	AD1928	875569	H82810	
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	Length	, ;	193	192	192	188	192	194	192	195	190	175	353	353	323	346	286	296	339	3033	1829	480	813	897	348	455	455	747	242	451	154	
•	A Query Match		100.0	90.4	89.8	84.3	82.9	81.0	80.5	80.4	80.1	75.4	10.6	10.5	10.4	10.3	10.2	9.8	9.8	9.5	9.4	9.3	9.3	9.3	9.5	9.5	9.5	9.5	9.5	9.1	9.0	
	Score		846	764.5	759.5	713	701	685.5	681	680.5	678	638	90	89	88	87.5	98	83	83	80.5	79.5	79	78.5		78	78	78	78	77.5	77	76.5	
	Result No.		-	7	m	4	ß	9	7	80	6	10	11	12	13	14	15	16	17	18	19	20	. 21	22	23	24	25	56	27	28	29	

mandelate racemase hypothetical prote conserved hypothet rts beta (AR305057 ATP-dependent heli RR2 protein saim thrombopoietin pre ribonucleoside-dip probable transport hypothetical prote GCNS protein hum VacB protein - hum VacB protein - hum VacB protein - hum VacB protein - hum VacB protein - hum	NADH2 dehydrogenas precorrin-6y c5,15
AE3465 875772 AB2922 C97696 D64738 B37994 JG4125 AH3625 AH3625 AH3625 AH3621 B5517 B9511 S71789 S71789 S7378	H69478 AF3341
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. 425 6425 6400 4200 824 228 336 1564 401 401 1089	379 401
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# ALIGNMENTS

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A;Cross-references: UNIPROT:P33709; EMBL:224081; NIU: 932001; RWHW, D; Boissel, J.
Blood 82, 1507-1516, 1993
A;Title: Brythropoietin structure-function relationships: High degree of sequence homol
A;Reference number: 146083; MUID:93372347; PMID:8364201
A;Reference number: 146083; MUID:93372347; PMID:8364201
A;Residues: translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 4-15,'','17-107,'P',109-194 < WEN>
A;Residues: 4-15,'','17-107,'P',109-194 < WEN>
A;Residues: 4-15,'','17-107,'P',109-194 < WEN>
A;Cross-references: GB:L10610; NID:9165876; PIDN:AAA31518.1; PID:9165877
C;Comment: Brythropoietin is produced by kidney or liver of adult mammals and by liver
C;Function:
A;Description: the primary inducer of erythrocyte formation
C;Superfamily: erythropoietin
C;Superfamily: erythropoietin
C;Superfamily: erythropoietin
C;Superfamily: erythropoietin
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.Стова-теferences; UNIPROT: P33709, EMBL: Z24681, NID:g395049; PIDN: CAA80848.1; PID:g395
Wen, D.; Boissel, J.
Lood 82, 1507-1516, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A,Cross-references: GB:M12930; NID:g193086; PIDN:AAA37570.1; PID:g387152
C,Comment: Erythropoietin is produced by kidney or liver of adult mammals and by liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: DNA
A; Residues: 1-192 < kNO-
A; Residues: 1-192 < kNO-
A; Cross-references: UNID-
B; Note: the authors translated the codon TTA for residue 12 as Phe, TTA for residue A; Note: the authors translated the codon TTA for residue 12 as Phe, TTA for residue 48, Note: the authors translated the codon TTA for residue 24, Note: the authors translated the codon TTA for residue 48, Note: TTA for residue 48, Note: TTA for residue 48, Note: TTA for residue 48, Note: TTA for residue 48, Note: TTA for residue 48, Note: TTA for residue 48, Note: TTA for residue 48, Note: TTA for residue 48, Note: TTA for residue 48, Note: TTA for residue 48, Note: TTA for residue 48, Note: TTA for residue 58, Note: TTA for residue 48, Note: TTA for residue 58, Note: TTA for r
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9
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C;Species: Mus musculus (house mouse)
C;Date: 25-Oct-1987 #sequence_revision 15-Nov-1996 #text_change 09-Jul-2004
C;Accession: A24902, A24901
R;Shoemaker, C.B.; Mitsock, L.D.
Mol. Cell. Biol. 6, 849-858, 1986
A;Title: Murine erythropoietin gene: cloning, expression, and human gene how
A;Reference number: A24902; MUID:87039105; PMID:3773894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28 APPRLICDSRVLERYILEAREAENATMGCAEGCSFSENITVPDTKVNFYAWKRMEVQQQA
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C; Reywords: erythropolesis; glycoprotein; predicted <SIG>
F;1-26/Domain: signal sequence #status predicted <MAT>
F;7-132/Product: erythropoletin #status predicted
F;31-187, 55-165/Disulfide bonds: #status predicted
F;50,64,109/Binding site: carbohydrate (Asn) (covalent) #status predicted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           148 LPDATPSAAPLRIFTVDALSKJFRIYSNFLRGKJTLYTGEACRRGD 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20; Indels
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81.9%; Pred. No. 3.9e-58;
ive 9; Mismatches 20
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C; Function:
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Best Local Similarity 81.9
Matches 136; Conservative
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A;Residues: 1-192 <NAG>
A;Cross-references: UNIPROT:P29676; GB:D10763; NID:g220735; PIDN:BAA01593.1; PID:g220736
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Blood 82, 1507-1516, 1993
A;Title: Errthropoietin structure-function relationships: High degree of sequence homold
A;Reference number: 146083; MUID:93372347; PMID:8364201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ď
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C;Comment: Erythropoietin is produced by kidney or liver of adult mammals and by liver
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                                                                VEVWQGLALLSEAVLRGQALLVNSSQPWEPLQLHVDKAVSGLRSLTTLLLRALGAQKBAIS 120
                                                                                                                                83 VEVWQGIALLISEAILRGQALLANSSQPSETLQLHVDKAVSSLRSLISLLRALGAQKEATS 142
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Species: Rattus norvegicus (Norway rat)
C;Date: 22-Nov-1993 #sequence_revision 15-Nov-1996 #text_change 09-Jul-2004
C;Accession: $28148; I62743
R;Nagao, M; Suga, H; Okano, M; Masuda, S; Narita, H; Ikura, K; Sasaki,
Biochim. Biophys. Acta 1171, 99-102, 1992
A;Title: Nucleotide sequence of rat erythropoietin.
A;Reference number: $28148; MUID:93042015; PMID:1420369
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C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Date: 16-Aug-1996 #sequence_revision 15-Nov-1996 #text_change 09-Jul-2004
C;Accession: 146401; 147077
R;Pu, P.; Evans, B.; Lim, G.B.; Moritz, K.; Wintour, E.M.
Mol. Cell. Endocrinol. 93, 107-116, 1993
A;Title: The sheep erythropoietin gene: molecular cloning and effect of hem
A;Reference number: 146401; MUID:93351736; PMID:8349021
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                                                                                                                                                                                                                                                                                                                                                                             PPDAASAAPLRTITADTFRKLFRVYSNFLRGKLKLYTGEACRTGD 165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       147 PPDATQAAPLRTLTADTFCKIFRVYSNPLRGKIKLYTGBACRRGD 191
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A,Molecule type: mRNA
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A; Residues: 4-192 <RES>
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Best Local

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C;Species: Yersinia pestis
C;Date: 02-Nov-2001 #sequence_revision 02-Nov-2001 #text_change 09-Jul-2004
C;Accession: AB0323
R;Parkhill, J; Wren, B.W.; Thomson, N.R.; Titball, R.W.; Holden, M.T.G.; Prentice, M.B deno-Tarraga, A.M.; Chillingworth, T.; Cronin, A.; Davies, R.M.; Davis, P.; Dougan, G.; Il, M.; Rutherford, K.; Simmonds, M.; Skelton, J.; Stevens, K.; Whitehead, S.; Barrell, Nature 413, 523-527, 2001
A;Title: Genome sequence of Yersinia pestis, the causative agent of plague.
A;Reference number: AB0001; MUID:21470413; PMID:11586360
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-33 < KURA.
A;Residues: 1-33 < KURA.
A;Residues: 1-33 < KURA.
A;Residues: 1-33 < KURA.
A;Residues: 1-33 < KURA.
A;Residues: 1-33 < KURA.
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AED0559
Solute binding receptor protein [imported] - Salmonella enterica subsp. enterica serova: C;Species: Salmonella enterica subsp. enterica serovar Typhi
S;Species: Salmonella enterica subsp. enterica serovar Typhi
A;Note: this species has also been called Salmonella typhi
C;Date: 09-Nov-2001 #sequence_revision 09-Nov-2001 #text_change 18-Nov-2002
C;Accession: AE0959
R;Parkhill, J; Dougan, G; James, K.D.; Thomson, N.R.; Pickard, D.; Wain, J.; Churcher th, T; Comnerton, P; Cronin, A.; Davis, P.; Davies, R.M.; Dowd, L.; White, N.; Parrar, S; Moule, S.; O'Gaora, P.
Nature 413, 848-852, 2001
A;Authors: Parry, C.; Quail, M.; Rutherford, K.; Simmonds, M.; Skelton, J.; Stevens, K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ribonucleoside-diphosphate reductase (BC 1.17.4.1) beta chain [imported] - Yersinia pes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    82 QDILGAVTLLLEGVMAARGQLGPTCLSSLLGQLSGQVRLLLGALQSL----LGTQ--- 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 VEVWOGLALLSEAVL--RGQALLVNSSQPWEPLQLHVDKAVSGLRSLTTLLRALGAQKEA 118
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                                                                                                                                                                                                                                                                                                                                                                                                                               1 APPRLICDSRVLERYLLEAKBAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA
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                                                                                                                                                                                                                                   10.5%; Score 89; DB 2; Length 353; 26.3%; Pred. No. 0.88; tive 20; Mismatches 75; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10.4%; Score 88; DB 2; Length 323; 25.2%; Pred. No. 0.99; tive 20; Mismatches 59; Indels
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A;Introns: 5,1; 47,3; 76,3; 132,3
C;Keywords: alternative splicing; cytokine; glycoprotein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  119 ISPPDAASAAPLRTITADTFRKLFRVYSNFLRGKLK 154
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C;Superfamily: ribonucleosi
C;Keywords: oxidoreductase
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Best Local Similarity
Matches 34; Conserva
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A;Accession: 180105
A;Status: translated from GB/EMBL/DDBJ
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C;Species: 24-May.1996 #text change 09-Jul-2004
C;Date: 24-May.1996 #sequence revision 24-May-1996 #text change 09-Jul-2004
C;Accession: ISS281; I80105; &45331; 848740; I38672; I52610
R;Poster, D.C.; Sprecher, C.A.; Grant, F.J.; Kramer, J.M.; Kuijper, J.L.; Holly, R.D.; W Proc. Natl. Acad. Sci. U.S.A. 91, 13022-13027, 1994
A;Title: Human thrombopoietin: gene structure, CDNA sequence, expression, and chromosoma A;Reference number: I59281; MUID:95108091; PMID:7809166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Residues: 1-353 <SAU>
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A; Cross-references: GB:L33410; NID:g506826; PIDN:AAA59857.1; PID:g506827
B; Sohma, X, Akahori, H.; Seki, N.; Hori, T.; Ogami, K.; Kato, T.; Shimada, Y.; Kawamura PEBS Lett. 353, 57-61, 1994
A; Title: Molecular cloning and chromosomal localization of the human thrombopoietin gene A; Reference number: S48740; MUID:95010765; PMID:7926023
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Bagenberger, J.; Hunt, P.; Li, Y.S.; Lu, H.S.; Martin, F.; Chang, M.S.;

Bartley, T.D.; Bogenberger, J.; Hunt, P.; Li, Y.S.; Lu, H.S.; Martin, F.; Chang, M.S.;

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A; Cf Sauvage, F.U.; Hass, P.E.; Spencer, S.D.; Malloy, B.E.; Gurney, A.L.; Spencer, S.A. D.V.; Eaton, D.L.
Nature 369, 533-538, 1994
A; Title: Stimulation of megakaryocytopolesis and thrombopolesis by the c-Mpl ligand.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               thrombopoietin precursor - human N;Alternate names: c-MPL ligand; megakaryocyte growth and development factor precursor C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Date: 24-May-1996 #text_change 09-Jul-2004 C;Date: 24-May-1996 #text_change 09-Jul-1004 C;Accession: 159281; 180105; 245331; S48740; 138672; 152810 R;Poster, D.C.; Sprecher, C.A.; Grant, P.J.; Kramer, J.M.; Kuijper, J.L.; Holly, R.D.;
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A;Reference number: S45331; MUID:94261202; PMID:8202154
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Gurney, A.L.; Kuang, W.J.; Xie, M.H.; Malloy, B.E.; Eaton, D.L.; de Sauvage, F.J.
Lood 85, 981-988, 1995
                                                                                                                                                                                  61 VEVWQGLALLSEAVL--RGQALLVNSSQPWEPLQLHVDKAVSGLRSLTTLLRALGAQKEA 118
                                                                                                                                                                                                                                                                           82 QDILGAVTLLLEGVMAARGQLGPTCLSSLLGQLSEQVRLLLGALQSL-----LGTQ--- 132
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                                                 24 APP--ACDLRVLSKLLRDSHVLHSKLSQCPEVHPLPTPVLLPAVDFSLGEWKTQMEETKA
APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA
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                                                                                                                                                                                                                                                                                                                                                                              119 ISPPDAASAAPLRTITADTFRKLFRVYSNFLRGKLK 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -LPPQG-----RTTAHKDPNAIFLSFQHLLRGKVR 161
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Residues: 1-353 <RE4>
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Status: preliminary
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OM protein - protein search, using sw model

Run on:

August 23, 2005, 13:52:31; Search time 178 Seconds (without alignments) 474.680 Million cell updates/sec

US-10-706-701-1 846 1 APPRLICDSRVLERYLLEAK......SNPLRGKLKLYTGBACRTGD 165 Title: Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

1612378 Total number of hits satisfying chosen parameters: 1612378 seqs, 512079187 residues Searched:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

UniProt\_03:\*
1: uniprot\_sprot:\*
2: uniprot\_trembl:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	P01588 homo sapien	P07865 macaca fasc	Q28513 macaca mula	н			Q6pwu5 canis famil	P48617 bos taurus	P07321 mus musculu		Q9gka2 oryctolagus		P49157 sus scrofa				8		-	-	_	Q8hz87 pongo pygma		Q8hz85 saguinus oe	Q6uam1 tetraodon n				gallu		P40225 homo sapien
Carvenios	GI.	EPO_HUMAN	BPO MACFA	EPO MACMU	Q867B1	EPO FELCA	EPO_RAT	QGPWUS		EPO MOUSE	EPO_SHEEP	Q9GKA2	Q9GKA3	RPO PIG	Q6H <u>8</u> S9	Q6H8T0	Q6H8T1	ОЭМУМВ	Q6H8T2	O8HZ88	Q8HZ89	BPO_CANFA	Q8HZ87	98ZH8Ō	Q8HZ85	Q6UAM1	Q6JV23	Q6JV22	Q9QV40	Q6IYB9	TPO_CANFA	TPO_HUMAN
	8 ;	-	П	-	7	7	г	7	Н	7	ч	N	0	-	7	~	7	7	(1	7	7	н	N	7	0	7	0	0	7	7	-	-
	Length	193	192	192	192	192	192	206	192	192	194	195	195	190	192	192	192	194	192	133	133	175	131	133	133	195	182	185	20	177	352	353
Query	Match	100.0	90.4	89.8	85.5	83.5	82.9	81.9	81.9	81.4	81.0	80.4	80.4	80.1	80.1	80.1	80.1	80.1	79.7	78.4	77.8	75.4	74.1	71.7	65.5	28.2	28.1	28.1	22.2	13.4	12.9	10.5
ć	Score	846	764.5	759.5	723	206	701	693	692.5	689	685.5	680.5	680.5	678	678	678	678	678	674	663	658	638	627	607	554	241	238	238	188	113	109	89
Result		7	63	М	4	ß	9	7	œ	σ	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	56	27	28	58	30	31

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## ALIGNMENTS

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                                                               VEVWOGLALLSEAVIRGOALIVNSSOPWEPLOLHVDKAVSGIRSLTTILLRALGAQKEAIS 120
                                                                                         88 VEVWQGLALLSEAVLRGQALLVNSSQPWEPLQLHVDKAVSGLRSLTTLLRALGAQKEAIS 147
        9
                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE-87052336; PubMed=2877922; DOI=10.1016/0378-1119(86)90183-6; Lin F.-K., Lin C.-H., Lai P.-H., Browne J.K., Egrie J.C., Smalling R., Fox G.M., Chen K.K., Castro M., Suggs S.; Monkey erythropoietin gene: cloning, expression and comparison with the human erythropoietin gene: cloning, expression and comparison with Gene 44:201-209(1986).

-I FUNCTION: Erythropoietin gene: ', the principal hormone involved in the regulation of erythrocyte differentiation and the maintenance of a physiological level of circulating erythrocyte mass.

-I SUBCELLULAR LOCATION: Secreted.
-I SUBCELLULAR LOCATION: Secreted.
-I TISSUB SPECIFICITY: Produced by kidney or liver of adult mammals and by liver of fetal or neonateal mammals.
-I SIMILARITY: Belongs to the EPO / TPO family.
similarity).
similarity).
similarity).
similarity).
                                                                                                                                                                                                                                                                                                                                                Macaca fascicularis (Crab eating macaque) (Cynomolgus monkey).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Cercopithecidae,
                                                                                                                          PIR, JO0173; JO0173.

HISSP: P01588; 1CN4.

InterPro; IPR0013013; EPO TPO.

InterPro; IPR001323; EPO TPO.

InterPro; IPR001321; Erythroptn.

PERM; PO1789; EPO TPO; 1.

PIRSP: PRSF001951; EPO; 1.

PRINTS; PRO0272; ERYTHROPTN.

PROSITE; PS00817; EPO TPO; 1.

Erythrocyte maturation; Glycoprotein; Hormone; Signal.

SIGNAL.
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Erythropoietin.

By similarity.

By similarity.

N-linked (GlCNAc. ...) (By N-linked (GlCNAc. ...) (By O-linked (GlCNAc. ...) (By W; EBA900F442AD4522 CRC64;
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01-AUG-1988 (Rel. 08, Last sequence update)
02-CCT-2004 (Rel. 45, Last annotation update)
Erythropoietin precursor.
                                                                                                                                                                                                                                          192
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Σ
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51
65
110
1152
21113 M
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                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9541;
                                                                                                                                                                                                                                       EPO MACFA
P07865;
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CARBOHYD
SEQUENCE
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                                                                                                                                                                                                               TISSUE-Kidney;

X MEDLINE-93372347; PubMed-8364201;

XA WED D., Boissel JD.P.R., Tracy T.E., Gruninger R.H., Mulcahy L.S.,

A Wen D., Boissel JD.P.R., Tracy T.E., Gruninger R.H., Mulcahy L.S.,

Czelusniak J., Goodman M., Bunn H.P.;

RI "Erythropoietin structure-function relationships: high degree of

RI "Erythropoietin structure-function relationships: high degree of

RI "Erythropoietin structure-function relationships: high degree of

RI "Erythropoietin structure-function relationships: high degree of

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RI "Erythropoietin structure-function relationships: high degree of

RI "Erythropoietin structure-function relationships: high degree of

RI "Erythropoietin 
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                                                                                                                                            28 APPRLICDSRVLERYLLEAKEAENVTWGCSESCSLNENITVPDTKVNFYAWKRMEVGQQA
                                                                                                             APPRLICOSRVLBRYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA
                                                             Gaps
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        DB 1; Length 192;
                                                                                                                                                                                                                                                                                                                         121 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKLKLYTGBACRTGD 165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mandaca mulatta (Rhesus macaque).
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleo
Mammalia, Butheria, Primates, Catarthini, Cercopithecidae,
                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; InterPro; InterPro; InterPro; IRBOULE, INTERPO158; EPO Tr.
PERM; PRO0728; EPO; T.
A PRINTS; PRO0272; ERYTHROPIN.
DR PROSITE; PS00817; EPO TPO; T.
W EXPLANCATE maturation; Glycoprotein; Hormon.
SIGNAL 28 192 By similarity.
34 187 By similarity.
60 By similarity.
61 By similarity.
61 By similarity.
62 By similarity.
63 By similarity.
64 By similarity.
65 By similarity.
65 By similarity.
66 By similarity.
67 By similarity.
67 By similarity.
68 By similarity.
69 By similarity.
60 By similarity.
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                                                             . 9
Query Match
90.4%; Score 764.5; DB 1
Best Local Similarity 91.5%; Pred. No. 9.6e-64;
Matches 151; Conservative 7; Mismatches 6
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01-NOV-1997 (Rel. 35, Last sequence update)
25-OCT-2004 (Rel. 45, Last annotation update)
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NCBI_TaxID=9544;
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               VEVWQGLALLSEAVLRGQALLVNSSQPWEPLQLHVDKAVSGLRSLTTLLRALGAQKEAIS 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE OF 4-192 FROM N.A.

SEQUENCE OF 4-192 FROM N.A.

SEQUENCE OF STRAIN=Spreague-Dawley; TISSUE=Kidney; MEDLINE=33372347; PubMed=8364201;
MEDLINE=33372347; PubMed=8364201;
Men D., Boissel J.P.R., Tracy T.E., Mulcahy L.S., Czelusniak J., Goodman M., Bunn H.F.;

Raythropodetin structure-function relationships: high degree of sequence homology among mammals.";

Blood 82:1507-1316(1993).

-I- FUNCTION: Expthropojectin is the principal hormone involved in the regulation of erythrocyte differentiation and the maintenance of a physiological level of circulating erythrocyte mass.

C. -I- SUBCELLULAR LOCATION: Secreted.

-I- TISSUE SPECIFICITY: Produced by kidney or liver of adult mammals and by liver of fetal or neonatal mammals.

C. -I- SIMILARITY: Belongs to the EPO / TPO family.
                                                                                                                                                                                                                                                                            Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
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Brythropoietin.

By similarity.

N-linked (GloNAc. . .) (By similarity).

N-linked (GloNAc. . .) (By similarity).
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STRAIN-Wistar; TISSUE-Kidney;
MEDLINE-93042015; Pubmed=1420369; DOI=10.1016/0167-4781(92)90146-Q;
Nagao M., Suga H., Okano M., Masuda S., Narita H., Ikura K.,
                                                                        121 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKLKLYTGEACRTGD 165
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RGD; 2559; EDO.
InterPro; IPR009079; 4 helix cytokine.
InterPro; IPR001313; EPO TPO.
InterPro; IPR001313; EPO TPO.
InterPro; IPR001313; EPO TPO.
INTERP; PRSP01951; EPO; 1.
PIRSP; PR00758; EPO TPO; 1.
PRNTS; PR00272; ERYTHROPTN.
PROSTIE; PR00177; EPO TPO; 1.
ERYTHROCYTE maturation; Glycoprotein; Hormone; Sig
                                                                                                                                                                                                                                                                                                                                                                                                                "Nucleotide sequence of rat erythropoietin.";
Biochim. Biophys. Acta 1171:99-102(1992).
                                                                                                                                                                                      01-APR-1993 (Rel. 25, Created)
01-APR-1993 (Rel. 25, Last sequence update)
25-OCT-2004 (Rel. 45, Last annotation update)
                                                                                                                                                             192 AA
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                                                                                                                                                                                                                                   Erythropoietin precursor.
                                                                                                                                                            STANDARD;
                                                                                                                                                                                                                                                                Rattus norvegicus (Rat)
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192
187
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                                                                                                                                      Gaps
109 N-linked (GlcNAc. . .) (By similarity). 21286 MW; 3EA632737E7D2443 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Carnivora; Pissipedia; Canidae; Canis.
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                                                                          Query Match 82.9%; Score 701; DB 1; Length 192; Best Local Similarity 82.4%; Pred. No. 9e-58; Matches 136; Conservative 13; Mismatches 16; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE-Kidney;
Souza D.S., Vicentim D.L., Costa F.F., Saad S.T.O.;
Submitted (MAR-2004) to the EMBL/GenBank/DDBJ databases.
EMBL, AYS72971; AAS77874.1; -.
RGO; GO:0005576; C:extracellular; IEA.
GO; GO:0005128; F:erythropoietin receptor binding; IEA.
GO; GO:0005179; F:hormone activity; IEA.
RINEEPRO; IPR009079; 4 helix_cytokine.
RINEEPRO; IPR009013; EFO_TPO.
RINEEPRO; IPR00313; EFO_TPO.
RINEEPRO; IPR00313; EFVENOPTN.
RPEam; PF00758; EPO_TPO; 1.
RRINES; PR00272; ERYTHROPTN.
RPROSITE; PS00217; EPO_TPO; 1.
SEQUENCE 206 AA; 22666 MW; IEBC64A02CE4F5B0 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        81.9%; Score 693; DB 2; Length 206; 81.2%; Pred. No. 5.5e-57; ive 13; Mismatches 18; Indels
                                                                                                                                                                                                                                                                                                                                                                                                   121 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKLKLYTGRACRTGD 165
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05-UUL-2004 (TrEMBLrel. 27, Last sequence update)
05-UUL-2004 (TrEMBLrel. 27, Last annotation update)
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Last annotation update)
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(Rel. 33, Last sequ
(Rel. 45, Last anno
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  109 1
192 AA;
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SEQUENCE FROM N.A.
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05-JUL-2004 (
05-JUL-2004 (
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01-FEB-1996
25-OCT-2004
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P48617;
     CARBOHYD
SEQUENCE
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EPO_BOVIN
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Pfam; PP00758; EPO TPO; 1.
PIRSF; PIRSP01951; EPO; 1.
PRINTS; PR00272; ERYTHROPTN.
PROSITE; PS00817; EPO TPO; 1.
SEQUENCE 195 AA; 21025 MW;
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194
189
60
51
65
110
108
                                                                                                           PIR; 146401; 146401.
ISSP; P01588; 1CN4.
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CARBOHYD
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Q9GKA2
   CCC DDR B DDR B DDR B DDR B DDR B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD B DD 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 VEVWQGLALLSEAVLRGQALLVNSSQPWEPLQLHVDKAVSGLRSLTTLLRALGAQKEAIS 120
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MEDLINE=93372347; PubMed=8364201;
MEDLINE=93372347; PubMed=8364201;
Men D., Boissel J.P.W., Tracy T.E., Gruninger R.H., Mulcahy L.S.,
Czelusniak J., Goodman M., Bunn H.F.;
"Erythropoietin structure-function relationships: high degree of
sequence homology among mammals.";
Blood 82:1507-1516 (1993).
-!-FUNCTION: Erythropoietin is the principal hormone involved in the
regulation of erythrocyte differentiation and the maintenance of
physiological level of circulating erythrocyte mass.
                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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-!- TISSUE SPECIFICITY: Produced by kidney or liver of adult mammals and by liver of fetal or neonatal mammals.
-!- SIMILARITY: Belongs to the BPO / TPO family.
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                                                                                                                                                 similarity).
similarity).
similarity).
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                                                                                                                                                                                                                                                                                                                                  Score 689; DB 1; Length 192;
Pred. No. 1.2e-56;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              147 PPDTTPPAPLRTLTVDTFCKLFRVYANFLRGKLKLYTGEVCRRGD 191
                                                                                                                                                                                                                                                                                                                                                                                                       Matches 132; Conservative 14; Mismatches 19; Indels
                                                                                                                                                 By similarity.
N-linked (GlcNAc. .) (By N-linked (GlcNAc. .) (By N-linked (GlcNAc. .) (By N-linked (GlcNAc. .) (By W, 65F94E214E0DEF2E CRC64;
Erythrocyte maturation; Glycoprotein; Hormone; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-FEB-1994 (Rel. 28, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
25-OCT-2004 (Rel. 45, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     194 AA
                                                                                                                                                                                                                                                                                                                                  81.4%; Score 689;
80.0%; Pred. No. 1
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                                                                                                                                                                                                                       109 N
21365 MW;
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                                                                                                                                                                                                                                                     192 AA;
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Best Local Similarity
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TISSUE=Kidney;
                                                                                                       DISULFID
CARBOHYD
CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EPO SHEEP
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SEQUENCE
                                      SIGNAL
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RPO_SHEEP
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61 VEVWOGILALLISEAVIRGOALLIVNSSOPWEPLOLHVDKAVSGLRSLITTLIRALGAQKEAIS 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oryctolagus cuniculus (Rabbit),
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Eutheria, Lagomorpha, Leporidae, Oryctolagus.
                                                                                                                                                                                                                                                                  (Potential)
                                                                                                                                                                                                                                                                                 .) (Potential)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            20; Indels
                                                                                                                                                                                                                                                                                                                          F -> L (in Ref. 2).
L -> P (in Ref. 2).
C025AAB0528131A9 CRC64;
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GO; GO:0005128; F:erythropoietin receptor binding; IEA.
GO; GO:0005179; F:hormone activity; IEA.
InterPro; IPR001979; 4 helix_cytokine.
InterPro; IPR001323; EPO_TPO.
InterPro; IPR001323; Erythroptn.
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                                                                                                                                                        Signal
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Last annotation update)
                                                                                                                                                                                                                By similarity.
By similarity.
By similarity.
N-linked (GlCNAc..
N-linked (GlCNAc..
InterPro; IPR009079; 4 helix cytokine.
InterPro; IPR001323; BFO TPO.
InterPro; IPR0030137; BFO TPO.
InterPro; IPR003013; BFO TPO.
Pfan; PFO0708; BFO TPO; 1.
PIRSF; PIRSF001951; BFO; 1.
PROSTTE; PR00817; BRYTROPTN.
BROSTTE; PR00817; BRYTROPTN.
Erythrocyte maturation; Glycoprotein; Hormone;
                                                                                                                                                                                                                                                                                                                                                                                                                                      .6e-56
                                                                                                                                                                                                                                                                                                                                                                                                                                                              9; Mismatches
                                                                                                                                                                            By similarity
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                                                                                                                                                                                                                                                                                                                                                                         21335 MW;
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                                                                                                                                                                                                     Shams I., Avivi A., Nevo E.; "Hypoxid strains and mole rat: Expression of "Hypoxid stress tolerance of the subterranean mole rat: Expression of erythropoietin and hypoxia-inducible factor-la."; Nucleic Acids Res. 0:0-0(2004).
                                                                                                                                                                                                                                                                                                             Shame I., Avivi A., Eviatar N.;
"Hypoxic stress tolerance of the blind subterranean mole rat:
"Hypoxic stress tolerance of the blind subterranean mole rat:
proc. Natl. Acad. Sci. U.S.A. 101:9698-9703 (2004).

EMBL; AJ715795; CAG29400.13.-.
GO; GO:0005128; F:erythropoietin receptor binding; IBA.
GO; GO:0005179; F:hormone activity; IEA.
InterPro; IPR003079; 4 helix cytokine.
InterPro; IPR003013; EPO TPO.
InterPro; IPR003013; Erythroptn.
PRINTS; PR00272; ERYTHROPIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                        Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Rodentia, Sciurognathi, Muridae, Spalacinae,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         erythropoietin.
; 72FCA94DE8C5AAB5 CRC64;
                                        Last sequence update)
Last annotation update)
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80.6%; Pred. No. 1.3e-55;
tive 8; Mismatches 24.
                                                                                                                                                                                                                                                                                   TISSUB=Liver;
PubMed=15210955; DOI=10.1073/pnas.0403540101;
192 AA
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PRT;
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                     05-JUL-2004 (TrEMBLrel. 27, 05-JUL-2004 (TrEMBLrel. 27, 05-JUL-2004 (TrEMBLrel. 27,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-JUL-2004 (TrEMBLrel. 27,
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Matches 133; Conservative
                                                                Erythropoietin precursor.
PRELIMINARY;
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                                                                                                                                              NCBI_TaxID=164323;
                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                           Spalax galili.
                                                                                                                                                                                      TISSUE=Liver;
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05-JUL-2004
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                                                                                Name=epo;
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Q6H8S9
                                                                                                                                    Spalax
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Q6H8T0
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completed: August 23, 2005, 13:55:39

Search completed: An Job time : 181 secs

Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi, Mammalia, Butheria, Rodentia, Sciurognathi, Muridae, Spalacinae,

Spalax judaei (Blind subterranean mole rat).

Erythropoietin precursor.

Name=epo;

Spalax.

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                                                                  Shams I., Avivi A., Nevo E.; "Hypoxic stress tolerance of the subterranean mole rat: Expression of erythropoietin and hypoxia-inducible factor-la."; Nucleic Acids Res. 0:0-0(2004).
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"Hypoxic stress tolerance of the blind subterranean mole rat:
"Hypoxic stress tolerance of the blind subterranean mole rat:
"Hypoxic stress tolerance of the blind subterranean mole rat:
Proc. Natl. Gerythropoietin and hypoxia.inducible factor I alpha.";
EMBL; AJ715794; CAG29399.1; -.
GO, GO:0005128; Castracellular; IEA.
GO, GO:0005128; Ferythropoietin receptor binding; IEA.
GO, GO:0005179; F.hormone activity; IEA.
InterPro; IPR003013; EPO TPO.
InterPro; IPR03013; EPYthroptn.
PRINTS; PR00272; ERYTHROPIN.
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; 72FCA94DE8C5AAB5 CRC64;
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GREEKAL INFORMATION:
APPLICANT: Burg, Josef
APPLICANT: Higer, Bernd
APPLICANT: Josel, Hans-Peter
TITLE OF INVENTION: ERYTHROPOIETIN CONJUGATES
FILE RFERENCE: 1098 nonprovisional
CURRENT APPLICATION NUMBER: US/09/604,871
CURRENT APPLICATION NUMBER: 60/15,454
FRIOR APPLICATION NUMBER: 60/147,452
FRIOR FILING DATE: 1999-08-05
FRIOR FILING DATE: 1999-07-02
FRIOR APPLICATION NUMBER: 60/147,423
FRIOR APPLICATION NUMBER: 60/142,243
FRIOR PILING DATE: 1999-07-02
NUMBER: OF SEQ ID NOS: 3
SOFTWARE: PALENTIN VEY: 2.1
SEQ ID NO 1
SEQ ID NO 1
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US-09-813-775C-32
US-09-552-265B-38
US-09-813-775C-38
US-09-813-775C-48
US-09-813-775C-48
US-09-813-775C-24
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; Sequence 1, Application US/09604938
; Patent No. 6583272
; GENERAL INFORMATION:
; APPLICANT: Ballon, Pascal
; TITLE OF INVENTION: ERYTHROPOIETIN CONJUGATES
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Patent No. 6340742
GENERAL INFORMATION:
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ORGANISM: Homo sapiens
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1 APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA 60
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100.0%; Pred. No. 1.5e-99;
iive 0; Mismatches 0;
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Patent No. 6608183
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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Best Local Similarity 100.0
Matches 165; Conservative
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; ORGANISM: Homo sapiens
US-09-462-941-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 121 PPDAASAAPLRTITADTFRKLFRVYSNPLRGKLKLYTGBACRTGD 165
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                                                                    APPLICANT: BLIEF, Josef
APPLICANT: Hilger, Bernd
APPLICANT: Hilger, Bernd
APPLICANT: Hilger, Bernd
APPLICANT: Josel, Hans-Peter
TITLE OF INVENTION: EXTITROPOIETIN CONJUGATES
FILE REFERENCE: 1098 nonprovisional
CURRENT APPLICATION NUMBER: US/09/604,871
CURRENT PILING DATE: 2000-06-28
PRIOR FILING DATE: 1999-08-30
PRIOR PRICATION NUMBER: 60/147,452
PRIOR FILING DATE: 1999-08-05
PRIOR FILING DATE: 1999-08-05
PRIOR FILING DATE: 1999-08-05
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TITLE OF INVENTION: ERYTHROPOIETIN CONJUGATES
FILE REPERRUCE: 1097 nonprovisional
CURRENT APPLICATION NUMBER: US/09/604,938
CURRENT PILING DATE: 2000-06-27
FRIOR APPLICATION NUMBER: 60/166,151
FRIOR FILING DATE: 1999-10-17
FRIOR FILING DATE: 1999-08-13
FRIOR FILING DATE: 1999-08-23
FRIOR APPLICATION NUMBER: 60/150,225
FRIOR APPLICATION NUMBER: 60/150,225
FRIOR APPLICATION NUMBER: 60/150,225
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FRIOR APPLICATION NUMBER: 60/150,225
FRIOR APPLICATION NUMBER: 60/150,225
FRIOR FILING DATE: 1999-00-23
FRIOR FILING DATE: 1999-00-23
FRIOR FILING DATE: 1999-00-23
FRIOR FILING DATE: 1999-00-23
FRIOR FILING DATE: 1999-07-02
FRIOR FILING DATE: 1999-07-02
Sequence 2, Application US/09604871
Patent No. 6340742
GENERAL INFORMATION:
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Patent No. 6583272
                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 3
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Homo sapiens
US-09-604-871-2
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LENGTH: 166
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GENERAL INFORMATION:
APPLICANT: Cox III, George N
APPLICANT: Bolder Blotechnology, Inc.
APPLICANT: Bolder Blotechnology, Inc.
TITLE OF INVENTION: Derivatives of Growth Hormone and Related Proteins
FILE REFERENCE: 4152-1-PUS
CURRENT APPLICATION NUMBER: US/09/462,941
CURRENT APPLICATION NUMBER: 60/052,516
PRIOR FILING DATE: 1997-07-14
NUMBER OF SEQ ID NOS: 41
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2
LENGTH: 166 ö APPLICANT: Brigham and Women's Hospital
APPLICANT: 75 Francis Street
APPLICANT: Boston, MA 02115
APPLICANT: Boston, MA 02115
APPLICANT: Bunn, H. Franklin
APPLICANT: Wen, Danyi
APPLICANT: Wen, Danyi
APPLICANT: Wen, Danyi
APPLICANT: Wen, Arrythropoietin Muteins With Enhanced
TITLE OF INVENTION: Activity
NUMBER OF SEQUENCES: 59
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox
STREET: 1100 New York Avenue, Suite 600
CITY: Washington Length 166; 121 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKLKLYTGEACRIGD 165 121 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKLKLYTGBACRTGD 165 121 PPDAASAAPLRIITADIFRKLFRVYSNFLRGKLKLYTGBACRTGD 165 121 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKLKLYTGEACRTGD 165 0; Indels

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88 VEVWQGLALLSEAVLRGQALLVNSSQPWEPLQLHVDKAVSGLRSLTTLLRALGAQKEAIS 147
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28 APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA 87
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                                                                                                                                                                                                                                                                                                       Sequence 4, Application US/09552265B

Patent No. 655343

GENERAL INFORMATION:
APPLICANT: Descavage, Frederick
APPLICANT: Henner, Dennis, J.
TITLE OF INVENTION: Polypeptides and nucleic acids encoding the same;
FILE REFERENCE: GENERY. 05/CP1
CURRENT APPLICATION NUMBER: US/09/552,265B
CURRENT FILING DATE: 2000-04-19
PRIOR APPLICATION NUMBER: US 09/307307
PRIOR PLING DATE: 1999-05-17
NUMBER OF SEQ ID NOS: 49

SOFTWARE: PASESEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: DeSauvage, Frederick
APPLICANT: Benner, Dennis, J.
TILLE OF INVENTION: No. 6831060el chimpanzee erythropoietin
TITLE OF INVENTION: No. 6831060el chimpanzee erythropoietin
TITLE OF INVENTION: No. 6831060el chimpanzee erythropoietin
TITLE OF INVENTION: NO. 6831060el chimpanzee erythropoietin
FILE REFERENCE: GENENT.057CP2
CURRENT PELICATION NUMBER: US/09/813,775C
CURRENT FILING DATE: 1999-05-07
PRIOR PELICATION NUMBER: US 09/307307
PRIOR PILING DATE: 1999-05-07
PRIOR FILING DATE: 2000-04-19
NUMBER OF SEQ ID NOS: 52
SUSTRARE: PASELSEQ for Windows Version 4.0
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                                                                                                                                             121 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKLKLYTGEACRTGD 165
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100.0%; Score 846; DB 4;
Best Local Similarity 100.0%; Pred. No. 1.9e-99;
Matches 165; Conservative 0; Mismatches 0;
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; ORGANISM: Homo sapiens
US-09-813-775C-4
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61 VEVWQGLALLSEAVLRGQALLVNSSQPWEPLQLHVDKAVSGLRSLTTLLRALGAQKEAIS 120
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                                                                                    1 APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA
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                                                Gaps
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  Length 193;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Jonathan Paul MURPHY
Anthony ATKINSON
TILLE OF INVENTION: Detection of Molecules in Samples
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSES: Pillabury Winthrop, L.L.P.
STREEF: 1100 New York Ave., N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         121 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKLKLYTGEACRTGD 165
                                          Indels
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100.0%; Score 846; DB 4;
100.0%; Pred. No. 1.9e-99;
iive 0; Mismatches 0;
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FILING DATE: No. 6680207ember 16, 1998
APPLICATION NUMBER: GB 9723955.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99.6%; Score 843; DB 4;
99.4%; Pred. No. 3.5e-99;
tive 1; Mismatches 0
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INFORMATION FOR SEQ ID NO: 8:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS WORTG
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/554,451
FILING DATE: 15-May-2000
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 8:
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Patent No. 6680207
GENERAL INFORMATION:
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LENGTH: 165 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A.
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: single
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Query Match
Best Local Similarity 100.0
Matches 165; Conservative
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Best Local Similarity 99.4°
Matches 164, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Published Applications AA:

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2: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

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3: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/US11_NEW_PUB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
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                                GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Maximum Match 100%
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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### ALIGNMENTS

%305-953-731-1 Sequence 1, Application US/09853731 Patent No. US20020037841A1 GENERAL INFORMATION: APPLICANT: Papadimitriou, Apollon TITLE OF INVENTION: Erythropoletin Composition FILE REFERENCE: 20619 US -1.rapb

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APPLICANI: CHEM, ALITE OF INVENTION: PROTEIN REMODELING METHODS AND PROTEINS/PEPTIDES PRODUCED BY THE TITLE OF INVENTION: METHODS

TITLE OF INVENTION: METHODS

FILE REPERENCE: 04083-301-503

CURRENT APPLICATION NUMBER: US 60/328,523

FRIOR PELING DATE: 2003-04-09

FRIOR PELING DATE: 2001-10-10

PRIOR APPLICATION NUMBER: US 60/344,692

PRIOR PILING DATE: 2000-10-19

PRIOR APPLICATION NUMBER: US 60/387,292

PRIOR PILING DATE: 2002-06-07

PRIOR PILING DATE: 2002-06-07

PRIOR PILING DATE: 2002-06-15

PRIOR PILING DATE: 2002-06-17

PRIOR PILING DATE: 2002-06-17

PRIOR PILING DATE: 2002-06-16

PRIOR PILING DATE: 2002-06-16

PRIOR PILING DATE: 2002-06-16

PRIOR PILING DATE: 2002-06-16

PRIOR PILING DATE: 2002-08-16

PRIOR PILING DATE: 2002-08-18

PRIOR PILING DATE: 2002-08-18

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; Pred. No. 1.4e-85;
0; Mismatches 0;
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Publication No. US20040063911A1
GENERAL INFORMATION:
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                     PRIOR FILING DATE: 2002-08-28
NUMBER OF SEQ ID NOS: 75
SOFTWARE: Patentin version 3.2
SEQ ID NO 73
LENGTH: 165
TYPE: PRT
ORGANISM: Homo sapiens
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Best Local Similarity 100.0%;
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APPLICANT: Deprees, Shawn

APPLICANT: Deprees, Shawn

APPLICANT: Applicant: Robert

APPLICANT: Bayer, Robert

APPLICANT: Bayer, Robert

APPLICANT: Hakes, David

APPLICANT: Chen, Xi

APPLICANT: Chen, Xi

APPLICANT: Chen, Xi

APPLICANT: Chen, Xi

APPLICANT: Chen, Xi

APPLICANT: Chen, Xi

APPLICANT: Chen, Xi

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APPLICANT: Chen, Xi

APPLICANT: Chen, Xi

APPLICANT: Chen, Xi

CURRENT APPLICANT: Chen, Xi

PRIOR PILING DATE: 2001-10-19

PRIOR PILING DATE: 2002-06-25

PRIOR PILING DATE: 2002-06-25

PRIOR PILING DATE: 2002-07-17

PRIOR APPLICATION NUMBER: US 60/396,594

PRIOR PILING DATE: 2002-07-17

PRIOR APPLICATION NUMBER: US 60/404,249

PRIOR PILING DATE: 2002-07-17

PRIOR APPLICATION NUMBER: US 60/407,527
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Sequence 1, Application US/10293551
Publication No. US20030120045A1
GENERAL INFORMATION:
APPLICANT: BAILON, PASCA1
TITLE OF INVENTION: ERYTHROPOIETIN CONJUGATES
FILE REPERSERE: 1097 nonproviational
CURRENT APPLICATION NUMBER: US/10/293,551
FRIOR APPLICATION NUMBER: US/09/604,938
PRIOR PILING DATE: 1909-11-17
PRIOR APPLICATION NUMBER: 60/166,151
PRIOR APPLICATION NUMBER: 60/166,151
PRIOR APPLICATION NUMBER: 60/151,548
PRIOR PILING DATE: 1999-08-13
PRIOR PILING DATE: 1999-08-13
PRIOR PILING DATE: 1999-08-13
PRIOR PILING DATE: 1999-08-23
PRIOR PILING DATE: 1999-08-23
PRIOR PILING DATE: 1999-08-23
PRIOR PILING DATE: 1999-08-23
PRIOR PILING DATE: 1999-08-23
PRIOR PILING DATE: 1999-07-02
SOFTWARE: PATENTING DATE: 1999-07-02
PRIOR PILING DATE: 1999-07-02
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Publication No. US20040043446A1
GRNERAL INFORMATION:
APPLICANT: Neose Technologies, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Homo sapiens
US-10-293-551-1
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APPLICANT: Chen, Xi
APPLICANT: Chen, Xi
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APPLICANT: Chen, Xi
APPLICANT: Chen, Xi
APPLICANT: Chen, Xi
APPLICANT: Chen, Xi
APPLICANT: Bowe, Caryn
TITLE OF INVENTION: FOLLICLE STIMULATING HORMONE: REMODELING AND GLYCOCONJUGATION OF TITLE OF INVENTION: FOLLICLE STIMULATING HORMONE: REMODELING AND GLYCOCONJUGATION OF TITLE OF INVENTION NUMBER: US/10/410,997
CURRENT APPLICATION NUMBER: US 60/328,523
PRIOR FILING DATE: 2001-10-19
PRIOR FILING DATE: 2001-10-19
PRIOR FILING DATE: 2002-06-25
PRIOR APPLICATION NUMBER: US 60/391,777
PRIOR FILING DATE: 2002-06-25
PRIOR FILING DATE: 2002-06-16
PRIOR FILING DATE: 2002-06-16
PRIOR FILING DATE: 2002-08-16
PRIOR FILING DATE: 2002-08-16
PRIOR FILING DATE: 2002-08-18
PRIOR FILING DATE: 2002-08-28
NUMBER OF SEQ ID NOS: 75
SEQ ID NO 73
TENCTH: 165
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US-10-411-012-73

Sequence 73, Application US/10411012

Sequence 73, Application Wol1041012

Sequence 73, Application Wol1041012

Sequence 73, Application Wol1040132640A1

GENERAL INFORMATION:

APPLICANT: Berees, Shawn

APPLICANT: Bayer, Robert

APPLICANT: Bayer, Robert

APPLICANT: Chen, Xi

APPLICANT: Bayer, Robert

APPLICANT: Bayer, Robert

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APPLICANT: Bayer, Robert

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APPLICANT: Bayer, Robert

APPLICANT: Bayer, Robert

APPLICANT: Chen, Xi

APPLICANT: Bayer, Robert

APPLICANT: Bayer, Robert

APPLICANT: Wolf and CAYCOPEGYLATION METHODS

TITLE OF INVENTION: METHODS

TITLE OF INVENTION: WETHODS

TITLE OF INVENTION NUMBER: US 60/328,523

PRIOR FILING DATE: 2001-10-19

PRIOR FILING DATE: 2001-10-19
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100.0%; Pred. No. 1.4e-85;
ative 0; Mismatches 0;
                                                                   DeFrees, Shawn
Zopf, David
Bayer, Robert
Hakes, David
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Best Local Similarity 100.
Matches 165; Conservative
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ORGANISM: Homo sapiens
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| Publication NO US20040115168A1
| Publication NO US20040115168A1
| Publication NO US20040115168A1
| APPLICANT: Depteres, Shawn
| APPLICANT: Depteres, Shawn
| APPLICANT: Copf, David
| APPLICANT: Bayer, Robert
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                                 1 APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA 60
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1 APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA
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; ORGANISM: Homo sapiens
US-10-410-930-73
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                                                                                                                                                                                                                                                                                                         1 APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    121 PPDAASAAPLRTITADTERKLFRVYSNFLRGKLKLYTGBACRIGD 165
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                                                                                                                                                                                                                                       Indels
                                                                                                                                                             100.0%; Score 846; DB 17;
100.0%; Pred. No. 1.4e-85;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION APPLICATION, Apollon, TITLE OF INVENTION: Erythropoletin Composition FILE REPRENCE: 20069 US CURRENT APPLICATION NUMBER: US/10/780,297 CURRENT FILING DATE: 2004-02-17 PRIOR APPLICATION NUMBER: US/09/853,731 PRIOR PILING DATE: 2000-05-11 PRIOR PLING DATE: 2000-05-11 PRIOR FILING DATE: 2000-05-15 NUMBER OF SEQ ID NOS: 2 SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application US/09853731
Patent No. US20020037841A1
GENERAL INFORMATION:
HAPLICANT: Bapadimitriou, Apollon
TITLE OF INVENTION:
CURRENT APPLICATION NUMBER: US/09/853,731
CURRENT APPLICATION NUMBER: EP/00110355.5
PRIOR FILING DATE: 2001-05-11
PRIOR FILING DATE: 2000-05-15
NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-780-297-1
; Sequence 1, Application US/10780297
; Publication No. US20040147431A1
; GENERAL INFORMATION:
                                                                                                                                                      Query Match
Best Local Similarity 100.0
Matches 165; Conservative
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Best Local Similarity 100.0
Matches 165; Conservative
                                                       ; ORGANISM: Homo sapiens
US-10-410-897-73
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US-10-780-297-1
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Publication No. US20050100982A1

GENERAL INFORMATION:

APPLICANT: DeFrees, Shawn

APPLICANT: DeFrees, Shawn

APPLICANT: DeFrees, Shawn

APPLICANT: DeFrees, Shawn

APPLICANT: DeFrees, Shawn

APPLICANT: Bayer, Robert

APPLICANT: Bayer, Robert

APPLICANT: Bayer, Robert

APPLICANT: Bayer, Robert

APPLICANT: Bayer, Robert

APPLICANT: Bowe, Caryn

ITILE OF INVENTION: FACTOR IX; REMODELING AND GLYCOCONJUGATION OF FACTOR IX

PRICANT: Define Dayer: 2003-04-09

PRIOR PILLING DATE: 2001-10-10

PRIOR PRILING DATE: 2001-10-10

PRIOR PILLING DATE: 2002-06-25

PRIOR PILLING DATE: 2002-06-26

PRIOR PILLING DATE: 2002-06-26

PRIOR PILLING DATE: 2002-07-17

PRIOR PILLING DATE: 2002-07-17

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PRIOR APPLICATION NUMBER: US 60/328,523
PRIOR FILING DATE: 2001-10-10
PRIOR PILING DATE: 2001-10-10
PRIOR PILING DATE: 2001-10-19
PRIOR PILING DATE: 2002-06-07
PRIOR PILING DATE: 2002-06-07
PRIOR PILING DATE: 2002-06-05
PRIOR PILING DATE: 2002-06-15
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PRIOR PILING DATE: 2002-08-18
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; ORGANISM: Homo sapiens
US-10-410-980-73
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LENGTH: 165
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61 VEVWQGLALLSBAVLRGQALLVNSSQPWEPLQLHVDKAVSGLRSLTTLLRALGAQKEAIS 120
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Publication No. US200301712841

GENERAL INFORMATION:

APPLICANT: Cox III, George N

TITLE OF INVENTION:

TITLE OF INVENTION DETIVATION OF US20101115

CURRENT APPLICATION NUMBER: US/10/298,148

CURRENT APPLICATION NUMBER: US/09/462,941

PRIOR APPLICATION NUMBER: 66/052,516

PRIOR APPLICATION NUMBER: 66/052,516

PRIOR APPLICATION NUMBER: 66/052,516

PRIOR PLING DATE: 2000-01-14

PRIOR PLING DATE: 1997-07-14

NUMBER OF SEQ ID NOS: 41

SOFTWARE: PatentIn Ver. 2.0
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                                                                     121 PPDAASAAPLRIITADTFRKLFRVYSNFLRGKLKLYTGEACRTGD 165
                                                                                               121 PPDAASAAPLRTITADTFRKUFRVYSNFLRGKLKLYTGEACRTGD 165
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Pred. No. 1.4e-85;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 227, Application US/10360101; Publication No. US20040009550A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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Best Local Similarity
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US-10-360-101-227
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                                                     Sequence 2, Application US/10400377

Publication No. US20030162949A1

GENERAL INFORMATION:

APPLICANT: Cox III, George N

APPLICANT: Cox III, George N

TITLE OF INVENTION: Derivatives of Growth Hormone and Related Proteins

TITLE OF INVENTION: Derivatives of Growth Hormone and Related Proteins

TITLE OF INVENTION: Derivatives of Growth Hormone and Related Proteins

CURRENT APPLICATION NUMBER: US/10/462,941

PRIOR APPLICATION NUMBER: US/09/462,941

PRIOR PILING DATE: 2000-01-14

PRIOR PILING DATE: 1997-07-14

PRIOR FILING DATE: 1997-07-14

SOFTWARE: PATENTING VOS: 41

SOFTWARE: PATENTIN VOS: 41

SOFTWARE: PATENTIN VOS: 41
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Publication No. US20030166865A1

GENERAL INFORMATION:

APPLICANT: Cox III, George N

APPLICANT: Bolder Biotechnology, Inc.

TITLE OF INVENTION: Derivatives of Growth Hormone and Related Proteins

FILE REFERENCE: 415.21-P07

CURRENT APPLICATION NUMBER: US/10/400,708

CURRENT PILING DATE: 2003-03-26

PRIOR PILING DATE: 2000-01-14

PRIOR FILING DATE: 1997-07-14

PRIOR FILING DATE: 1997-07-14

NUMBER OF SEQ ID NOS: 41

SOFTWARE: PatentIN Ver. 2.0

SEQ ID NO 2

LENGTH: 166
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Best Local Similarity 100.0
Matches 165; Conservative
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ORGANISM: Homo sapiens
                                            US-10-400-377-2
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US-10-400-708-2
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NUMBER OF SEQ ID NOS: 2036
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 133
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Sequence 2, Application US/10866540

Publication No. US20040230040A1

GENERAL INFORMATION:

APPLICANT: Cox III, George N
                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.
Matches 165; Conservative
                                                                                                                  TYPE: PRT
ORGANISM: Homo Sapiens
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; Bedication No. US20040180386A1
; GENERAL INFORMATION:
APPLICANT: Cartr, Graham
; APPLICANT: Carter, Graham
; APPLICANT: Carter, Graham
; APPLICANT: Hamilton, Anita
TITLE OF INVENTION: EPITOPES AND USE FOR PREPARING MOLECULES WITH REDUCED
; TITLE OF INVENTION: EPITOPES AND USE FOR PREPARING MOLECULES WITH REDUCED
; TITLE OF INVENTION: METHOD FOR IDENTIFICATION OF T-CELL
TITLE OF INVENTION: METHOD FOR IDENTIFICATION OF T-CELL
TITLE OF INVENTION: METHOD FOR IDENTIFICATION OF T-CELL
TITLE OF INVENTION: METHOD FOR IDENTIFICATION OF T-CELL
TITLE OF INVENTION: METHOD FOR IDENTIFICATION NUMBER: 01103954.2
FRIOR APPLICATION NUMBER: 01105536.4
FRIOR PILING DATE: 2001-03-15
FRIOR PILING DATE: 2001-03-15
FRIOR PILING DATE: 2001-03-15
FRIOR PILING DATE: 2001-03-15
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FRIOR PILING DATE: 2001-03-15
FRIOR PILING DATE: 2001-03-15
FRIOR PILING DATE: 2001-03-20
FRIOR PILING DATE: 2001-03-20
FRIOR FILING DATE: 2001-03-20
                                                                                                                           Sequence 2, Application US/10714149

Fublication No. US20040175800A1

GENERAL INFORMATION:

APPLICANT: Cox III, George N

APPLICANT: Cox III, George N

TITLE OF INVENTION Derivatives of Growth Hormone and Related Proteins

TITLE OF INVENTION NUMBER: US/10/774,149

CURRENT FILING DATE: 2004-02-05

FRIOR APPLICATION NUMBER: US/10/400,377

FRIOR APPLICATION NUMBER: US/09/462,941

FRIOR PILING DATE: 2000-01-14

FRIOR PILING DATE: 2000-01-14

FRIOR FILING DATE: 1997-07-14

NUMBER OF SEQ ID NOS: 41

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 2

LENGTH: 166
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  121 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKLKLYTGEACRTGD 165
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| Publication No. US20040214287A1
| GENERAL INFORMATION:
| APPLICANT: Cox III, George N
| APPLICANT: Cox III, George N
| TITLE OF INVENTION: Derivatives of Growth Hormone and Related Proteins
| TITLE OF INVENTION: Derivatives of Growth Hormone and Related Proteins
| TITLE OF INVENTION: Derivatives of Growth Hormone and Related Proteins
| TILE REPRENCE: 4152-1-PUS
| CURRENT FILING DATE: 2004-02-05
| PRIOR APPLICATION NUMBER: US/10/400,377
| PRIOR PILING DATE: 2003-03-26
| PRIOR FILING DATE: 2000-01-14
| PRIOR FILING DATE: 2000-01-14
| PRIOR FILING DATE: 2000-01-14
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| PRIOR FILING DATE: 2000-01-14 9 1 APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNPYAWKRMEVGQQA 60 1 APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRAEVGQQA 1 APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA 1 APPRLICDSRVLERYLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA Gaps Gaps ö ô Length 166; Length 166; 121 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKLKLYTGEACRTGD 165 121 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKCKLYCYTGEACRTGD 165 Indels Indels 100.0%; Score 846; DB 16; 100.0%; Pred. No. 1.4e-85; cive 0; Mismatches 0; Query Match 100.0%; Score 846; DB 16; Best Local Similarity 100.0%; Pred. No. 1.4e-85; Matches 165; Conservative 0; Mismatches 0;

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TYPE: PRT
ORGANISM: CHO/dhfr-
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Publication No. US20050107591A1

GENERAL INFORMATION:

APPLICANT: Cox III, George N

APPLICANT: Cox III, George N

TITLE OF INVENTION: Derivatives of Growth Hormone and Related Proteins

TITLE OF INVENTION: Derivatives of Growth Hormone and Related Proteins

TITLE OF INVENTION: Derivatives of Growth Hormone and Related Proteins

TITLE OF INVENTION: Derivatives of Growth Hormone and Related Proteins

TITLE OF INVENTION NUMBER: US/10/400,377

PRIOR APPLICATION NUMBER: US/09/462,941

PRIOR APPLICATION NUMBER: US/09/462,941

PRIOR APPLICATION NUMBER: 60/652,516

PRIOR PILING DATE: 2000-01-14

PRIOR PILING DATE: 1997-07-14

NUMBER OF SEQ ID NOS: 41

SOFTWARE: PATENTIN VET: 2.00

SEQ ID NOS: 41
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Best Local Similarity 100.0%; Pred. No. 1.4e-85;
Matches 165; Conservative 0; Mismatches 0;
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Pred. No. 1.4e-85;
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100.0%; Pred. No. ...
0; Mismatches
CURRENT FILING DATE: 2004-06-10
PRIOR APPLICATION NUMBER: US/10/400,377
PRIOR PILING DATE: 2003-03-26
PRIOR PILING DATE: 2003-03-26
PRIOR PILING DATE: 2000-01-14
PRIOR PILING DATE: 2000-01-14
PRIOR PILING DATE: 1997-07-14
NUMBER OF SEQ ID NOS: 41
SEQ ID NO 2
LENGTH: 166
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                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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US-10-773-530-2
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61 VEVWQGLALLSEAVLRGQALLVNSSQPWEPLQLHVDKAVSGLRSLTTLLRALGAQKEAIS 120 61 VEVWQGLALLSEAVLRGQALLVNSSQPWEPLQLHVDKAVSGLRSLTTLLRALGAQKEAIS 120 1 APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA 60 1 APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA Gaps ö Length 166; 121 PPDAASAAPLRTITADTFRKLFRVXSNFLRGKIKLYTGEACRIGD 165 121 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKLKLYTGEACRTGD 165 Indels 100.0%; Score 846; DB 18; 100.0%; Pred. No. 1.4e-85; ative 0; Mismatches 0; Sequence 2, Application US/10780297
Publication No. US20040147431A1
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Erythropoietin Composition
FILE REFRENCE: 200619 US
CURRENT APPLICATION NUMBER: US/10/780,297
CURRENT PILING DATE: 2004-02-17
PRIOR FILING DATE: 2001-05-11
PRIOR FILING DATE: 2001-05-11
PRIOR FILING DATE: 2000-05-15
PRIOR FILING DATE: 2000-05-15
PRIOR PRILING DATE: 2000-05-15
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin version 3.0
SEQ ID NO 2 Query Match Best Local Similarity 100.0 Matches 165; Conservative TYPE: PRT ORGANISM: Homo sapiens LENGTH: 166 US-10-780-297-2

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Sequence 4, Application US/10014363; Publication No. US20020115833A1; GENERAL INFORMATION: US-10-014-363-4

APPLICANT: Burg, Josef
APPLICANT: Burg, Josef
APPLICANT: Franze, Reinhard
APPLICANT: Hilger, Bernd
APPLICANT: Tischer, Wilhelm
APPLICANT: Tischer, Wilhelm
APPLICANT: Wazny, Manfred
TITLE OF INVENTION: Erythropoietin Conjugates
FILE REFERENCE: Case 20805
CURRENT APPLICATION NUMBER: US/10/014,363
CURRENT APPLICATION NUMBER: 2001-12-11
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin version 3.1 ENGTH: 169 SEQ ID NO 4

Gaps ò Length 169; ; Score 846; DB 13; Length 1; Pred. No. 1.4e-85; 0; Mismatches 0; Indels Query Match 100.0%; Best Local Similarity 100.0%; Matches 165; Conservative 0 4 APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA 63

1 APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA

61 VEVWQGLALLSEAVLRGQALLVNSSQPWEPLQLHVDKAVSGLRSLTTLLRALGAQKEAIS 120

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100.0%; Score 846; DB 16; Length 193; 100.0%; Pred. No. 1.7e-85; ive 0; Mismatches 0; Indels 0
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                                                                         121 PPDAASAAPLRIITADIFRKLFRVYSNFLRGKLKLYTGBACRTGD 165
                                                                                                 148 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKLKLYTGEACRTGD 192
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Publication No. US20040122216A1
GENERAL INFORMATION:
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; Publication No. US20040122216Al
; GENERAL INFORMATION:
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Best Local Similarity 100.0
Matches 165; Conservative
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Kallunki, P.
Christensen,
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Pedersen, L.
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Gerwien, J.
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APPLICANT: Pedersen, J
APPLICANT: Gervien, J.
APPLICANT: Bay, K.
APPLICANT: Pedersen, L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Artificial
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APPLICANT:
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APPLICANT: Sager, T.
APPLICANT: Brines, M.
APPLICANT: Brines, M.
APPLICANT: Brines, M.
APPLICANT: Cerami, A.
APPLICANT: Cerami, A.
TITLE OF INVENTION: RECOMBINANT TISSUE PROTECTIVE CYTOKINES AND ENCODING NUCLEIC
TITLE OF INVENTION: RESPONSIVE CELLS, TISSUES AND ORGANS
TITLE OF INVENTION: RESPONSIVE CELLS, TISSUES AND ORGANS
TILE OF INVENTION: RESPONSIVE CELLS, TISSUES AND ORGANS
TILE OF INVENTION: RESPONSIVE CELLS, TISSUES AND ORGANS
TILE OF INVENTION: BASEONSIVE CELLS, TISSUES AND ORGANS
TILE REFERENCE: 10165-022-999
CURRENT FILING DATE: 2003-07-01
PRIOR APPLICATION NUMBER: 60/392,455
PRIOR FILING DATE: 2002-07-01
PRIOR FILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-07-03
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                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                   Query Match 100.0%; Score 846; DB 14; Best Local Similarity 100.0%; Pred. No. 1.7e-85; Matches 165; Conservative 0; Mismatches 0;
PRIOR FILING DATE: 2002-01-04
PRIOR PELING NUMBER: US 60/358,598
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 22
SOFTWARE: Patentin version 3.1
SEQ ID NO 2
LENGTH: 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 10, Application US/10612665
Publication No. US20040122216A1
GENERAL INFORMATION:
APPLICANT: Nielsen, J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin version 3.2
SEQ ID NO 10
LENGTH: 193
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Geist, M.
Kallunki, P.
Christensen, S.
Sager, T.
Brines, M.
Cerami, A.
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                                                                                                                                                    TYPE: PRT
ORGANISM: Homo sapiens
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Pedersen, L.
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; ORGANISM: Homo sapiens
US-10-612-665-10
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APPLICANT: Sager, T.
APPLICANT: Sager, T.
APPLICANT: Brines, M.
APPLICANT: Cerami, A.
APPLICANT: Cerami, A.
APPLICANT: Cerami, A.
APPLICANT: Cerami, A.
APPLICANT: Cerami, C.
TITLE OF INVENTION: RECOMBINANT TISSUE PROTECTIVE CYTOKINES AND ENCODING NUCLEIC
TITLE OF INVENTION: RESPONSIVE CELLS, TISSUES AND ORGANS
TITLE OF INVENTION: RESPONSIVE CELLS, TISSUES AND ORGANS
TILLE OF INVENTION: RESPONSIVE CELLS, TISSUES AND ORGANS
CURRENT APPLICATION NUMBER: US/10/612,665
CURRENT FILING DATE: 2002-07-01
PRIOR PELING DATE: 2002-07-01
PRIOR PELING DATE: 2002-07-03
PRIOR PELING DATE: 2002-07-03
PRIOR FILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-07-03
PRIOR PELING DATE: 2002-07-03
PRIOR PELING DATE: 2002-07-03
PRIOR PELING DATE: 2002-07-03 61 VEVWQGLALLISEAVIRGOALLIVNSSQPWEPLQLHVDKAVSGLRSLTTLLRALGAQKEAIS 120 28 APPRLICDSRVLERYLLEAKBAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA 88 VEVWQGLALLSEAVLRGQALLVNSSQPWEPLQLHVDKAVSGLRSLTTLLRALGAQKEAIS 1 APPRLICDSRVLERYLLEAKGAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA

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Sequence 1, Application US/11021516
; Sequence 1, Application US/11021516
; Bublication No. US20650170457A1
; GENERAL INFORMATION:
; APPLICANT: CHICCOX. Inc.
; APPLICANT: Cunningham, Mark
; APPLICANT: Mills, Juliane
; APPLICANT: Mills, Juliane
; TITLE OF INVENTION: NOVEL RECOMBINANT PROTEINS WITH N-TERMINAL FREE THIOL
; FILE REFERENCE: CEN 5046
; CURRENT PAPLICATION NUMBER: 2004-12-23
; FRIOR APPLICATION NUMBER: 2004-12-23
; PRIOR PILING DATE: 2003-12-31
; NUMBER OF SEQ ID NOS: 20
; SOFTWARE: Patentin Version 3:3
; SEQ ID NOS: 20
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                                                                                   88 VEVWQGLALLSBAVLRGQALLVNSSQPWEPLQLHVDKAVSGLRSLTTLLRALGAQKKAIS 147
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Publication No. US20050170457A1
GENERAL INFORMATION:
APPLICANT: Centocor, Inc.
APPLICANT: Cuningham, Mark
APPLICANT: Pool, Chadler
APPLICANT: Pool, Chadler
APPLICANT: Pool, Chadler
APPLICANT: Sol, Solder
TITLE OF INVENTION: NOVEL RECOMBINANT PROTEINS WITH N-TERMINAL FREE THIOL
FILE REFERENCE: CEN 5046
CURRENT APPLICATION NUMBER: US/11/021,516
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                                                                                                                                                  121 PPDAASAAPLRIITADTFRKLFRVYSNFLRGKLKLYTGEACRTGD 165
                                                                                                                                                                            148 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKCKCYTGEACRTGD 192
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100.0%; Score 846; DB 20;
Best Local Similarity 100.0%; Pred. No. 1.7e-85;
Matches 165; Conservative 0; Mismatches 0;
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; OTHER INFORMATION: TRUNCATION, desarg
US-11-021-516-1
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LOCATION: (28)..(193)
FEATURE:
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APPLICANT: Fiordaliso, F.
APPLICANT: Fiordaliso, F.
APPLICANT: Goldiso, F.
APPLICANT: Gido, G.
TITLE OF INVENTION: TISSUE PROTECTIVE CYTOKINE RECEPTOR COMPLEX AND ASSAYS FOR IDENTI
TITLE OF INVENTION: TISSUE PROTECTIVE COMPOUNDS
FILE REPRENCE: 10165-027-999
CURRENT APPLICATION NUMBER: US/10/676,694
CURRENT FILING DATE: 2003-09-30
FRIOR APPLICATION NUMBER: 60/465,891
FRIOR PLILING DATE: 2003-04-25
NUMBER OF SEQ ID NOS: 212
SOOTWARE: Patentin version 3.2
SEQ ID NO 112
LENGTH. 193
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TITLE OF INVENTION: HIGH LEVEL EXPRESSION OF RECOMBINANT HUMAN ERYTHROPOIETIN
TITLE OF INVENTION: HAVING
TITLE OF INVENTION: HAVING
TITLE OF INVENTION: A MODIFIED 5'-UTR
PILE REPERENCE: 27179
CURRENT APPLICATION NUMBER: US/10/759,031
CURRENT PILING DATE: 2004-01-20
NUMBER OF SEQ ID NOS: 13
SEQ ID NO 10
LENGTH: 193
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      121 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKLKLYTGEACRTGD 165
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Publication No. US20050158822A1
GENERAL INFORMATION:
Publication No. US20040214236A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               , ORGANISM: Homo sapiens
US-10-759-031-10
                                                APPLICANT: Nielsen, M.
                                                                                                                                                                         Brines, M.
Cerami, A.
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US-10-759-031-10
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JUNEAU TO APPLICATION US/10622108

Publication No. US20040063912A1

SEQUENCE 10, Application US/10622108

Publication No. US20040063912A1

SEQUENCE INFORMATION:

APPLICANT: Blumberg, Mayne I.

APPLICANT: Bitonti, Alan J.

TILE REFERENCE: S01383.70011.US

CURRENT APPLICATION NUMBER: US/10/622,108

CURRENT FILING DATE: 2003-07-17

PRIOR PELING DATE: 2003-07-17

PRIOR PELING DATE: 2003-05-09

PRIOR PELING DATE: 2002-07-03

PRIOR PELING DATE: 2002-07-03

PRIOR PELING DATE: 2002-07-03

PRIOR PELING DATE: 2002-07-03

PRIOR PELING DATE: 2002-03-15

NUMBER OF SEQ ID NOS: 40

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100.0%; Pred. No. 5.2e-85;
ive 0; Mismatches 0;
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          FILE REFERENCE: S01383.70010.US
CURRENT APPLICATION NUMBER: US/10/435,608
CURRENT FILING DATE: 2003-05-09
PRIOR APPLICATION NUMBER: PCT/US02/21335
PRIOR FILING DATE: 2002-07-03
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin version 3.1
SEQ ID NO 10
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ORGANISM: Homo sapiens
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US-10-622-108-10
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Publication No. US20030235536A1
Publication No. US20030235536A1
APPLICANT: Blumberg, Richard S.
APPLICANT: Lencer, Wayne I.
APPLICANT: Sinistecr, Neal B.
APPLICANT: Bitonti, Alan J.
TITLE OF INVENTION: CENTRAL AIRWAY ADMINISTRATION FOR SYSTEMIC DELIVERY OF THERAPEUTI
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APPLICANT: MYUNG-EOK, CH
APPLICANT: MYUNG-EOK, CH
APPLICANT: JI-SOOK, PARK
APPLICANT: JI-SOOK, PARK
APPLICANT: JI-SOOK, PARK
TITLE OF INVENTION: FUSION PROTEIN HAVING ENHANCED IN VIVO ACTIVITY OF
TITLE OF INVENTION: ERYTHROPIETIN
FILE REFERENCE: 58105 (71970)
CURRENT PELLING DATE: 2002-08-29
PRIOR PAPLICATION NUMBER: 2001-74975
PRIOR PLILING DATE: 2001-11-29
NUMBER OF SEO ID NOS: 18
SOFTWARE: PATENTIN VET: 2.1
                                                                                                                                                                                                                     28 APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA
                                                                                                                                                        1 APPRLICDSRVLERYLLEAKEAENITTGCAEHCSLNENITVPDTKVNFYAWKRMEVGQQA
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                     Length 220;
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100.0%; Score 846; DB 14; Length 370;
Best Local Similarity 100.0%; Pred. No. 4.2e-85;
Matches 165; Conservative 0; Mismatches 0; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                121 PPDAASAAPLRTITADTFRKLFRVYSNFLRGKLKLYTGEACRTGD 165
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                                                                                   Indels
              100.0%; Score 846; DB 14;
100.0%; Pred. No. 2.1e-85;
ive 0; Mismatches 0;
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LENGTH: 370
TYPE: PRT
ORGANISM: Artificial Sequence
Query Match
Best Local Similarity 100.0
Matches 165; Conservative
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US-10-435-608-10
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APPLICANT: Sun, Lee-Hwei K
APPLICANT: Sun, Lee-Hwei K
APPLICANT: Sun, Bill N
APPLICANT: Sun, Bill N
APPLICANT: Sun, Cecily R
TITLE OF INVENTION: Fc fusion proteins of human erythropoietin with increased biolog PILE REPERENCE: 02SUN2001
CURRENT APPLICATION NUMBER: US/09/932, 812
CURRENT APPLICATION NUMBER: US/09/932, 812
SUFFMENT FILING DATE: 2001-10-30
NUMBER OF SEQ ID NOS: 22
SOFTWARE: Patentin version 3.1
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APPLICANT: Sun, Lee-Hwei K
APPLICANT: Sun, Bill N
TITLE OF INVENTION: Activities
TITLE OF INVENTION: activities
TITLE OF PRERENCE: 0250/02001-A
CURRENT APPLICATION NUMBER: US/10/761,593A
CURRENT FILING DATE: 2004-01-21
                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: HuEPO-L-vPc gamma2 with a 27-amino acid leader peptide (Pigure
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 846; DB 10; 100.0%; Pred. No. 5.3e-85;
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     Sequence 18, Application US/09932812
Publication No. US20030082749A1
GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: 09/932812
PRIOR FILING DATE: 2001-08-17
NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin version 3.2
                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 165; Conservative
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Sequence 22, Application US/11017185

Sequence 22, Application US/11017185

Sequence 22, Application US/11017185

Sequence 22, Application US/11017185

Sequence 22, Application US/11017185

Sequence 22, Application No. USC005014264

APPLICANT: Sun, Lee-Hwei K

APPLICANT: Sun, Cecily R

TITLE OF INVENTION: Ectivities

TITLE OF INVENTION: Cecily R

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TITLE OF INVENTION: Cecily R

FILE REPRENCE: 025UNZ001D2

CURRENT APPLICATION NUMBER: US/11/017,185

CURRENT FILING DATE: 2001-08-17

NUMBER OF SEQ ID NOS: 28

SOFTWARE: Patentin version 3.1

SEQ ID NO 22

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LEASTH: 435
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OTHER INFORMATION: )
                                                                                                                                                                             OTHER INFORMATION: HuEPO-L-vFc gammal with a 27-amino acid leader peptide (Figure OTHER INFORMATION: 2C)
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100.0%; Score 846; DB 20; Length 435;
Best Local Similarity 100.0%; Pred. No. 5.3e-85;
Matches 165; Conservative 0; Mismatches 0; Indels 0
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; Pred. No. 5.3e-85;
0; Mismatches 0;
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Best Local Similarity 100.0%;
Matches 165; Conservative 0
                                                                                                   TYPE: PRT ORGANISM: Artificial Sequence
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NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin version
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US-09-932-812-18
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                                                 SEQ ID NO 22
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OTHER INFORMATION: HuEPO-L-vFc gamma4 with a 27-amino acid leader peptide (Figure CTHER INFORMATION: 2B)
US-10-761-593A-20
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Sequence 20, Application US/20050124045A1

GENERAL INFORMATION:

APPLICANT: Sun, Lee-Hwei K

APPLICANT: Sun, Cecily R

TITLE OF INVENTION: Fc fusion proteins of human erythropoietin with increased

TITLE OF INVENTION: biological activities

FILE REFERENCE: 0.25UN2004D1

CURRENT APPLICATION NUMBER: US/11/016,518A

CURRENT FILING DATE: 2004-12-17

PRIOR PILING DATE: 2001-08-17

NUMBER OF SEQ ID NOS: 28

SOFTWARE: PatentIn version 3.2

SEQ ID NO 20

LENGTH: 437
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US-11-016-518A-20
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100.0%; Pred. No. 5.3e-85;
tive 0; Mismatches 0;
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CURRENT FILING DATE: 2004-01-21
PRIOR APPLICATION NUMBER: 09/932812
PRIOR FILING DATE: 2001-08-17
NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin version 3.2
SEQ ID NO 20
LENGTH: 437
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Best Local Similarity 100.0
Matches 165; Conservative
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Sequence 20, Application US/11017185

Publication No. US20050142642A1

GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TAPLICANT: Sun, Lee-Hwei K
APPLICANT: Sun, Cecily R
TITLE OF INVENTION: Activities
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TITLE OF INVENTION WHABER: US/11/017,185
CURRENT APPLICATION NUMBER: US/11/017,185
CURRENT FILING DATE: 2004-12-17
NUMBER OF SEQ ID NOS: 28
SOFUMARE: PATENTING PATE: 2010-08-17
NUMBER OF SEQ ID NOS: 28
SOFUMARE: PATENTING PATE: 2010-08-17
SEQ ID NOS: 28
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ORGANISM: Artificial Sequence
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